

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: July 20, 2004, 09:29:26 ; Search time 14 Seconds
(without alignments)

Sequence: 1 PQRKTKRNTNRRPQDKFPG 20

Title: US-10-044-995-2

Perfect score: 110

Scoring table: BL03M62

Gapext: 0.5

Searched:

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents AA:*

1: /cgcn2_6/.ptodata/2/iaa/5A_COMB.pep:*

2: /cgcn2_6/.ptodata/2/iaa/5B_COMB.pep:*

3: /cgcn2_6/.ptodata/2/iaa/6A_COMB.pep:*

4: /cgcn2_6/.ptodata/2/iaa/6B_COMB.pep:*

5: /cgcn2_6/.ptodata/2/iaa/PCUTS_COMB.pep:*

6: /cgcn2_6/.ptodata/2/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	Match	Length	DB	ID	Description
1	110	100.0	20	2	US-08-466-975A-2	Sequence 2, Appli	
2	110	100.0	20	2	US-08-391-671A-2	Sequence 2, Appli	
3	110	100.0	20	3	US-08-467-902A-2	Sequence 2, Appli	
4	110	100.0	20	3	US-09-275-265-2	Sequence 2, Appli	
5	110	100.0	20	4	US-09-941-611-2	Sequence 2, Appli	
6	110	100.0	20	4	US-09-790-497A-38	Sequence 38, Appli	
7	110	100.0	22	2	US-08-146-028-38	Sequence 38, Appli	
8	110	100.0	22	2	US-08-146-028-118	Sequence 118, Appli	
9	110	100.0	22	2	US-08-146-028-134	Sequence 114, Appli	
10	110	100.0	22	3	US-08-723-425A-38	Sequence 38, Appli	
11	110	100.0	22	3	US-08-723-425A-118	Sequence 118, Appli	
12	110	100.0	22	3	US-08-723-425A-134	Sequence 114, Appli	
13	110	100.0	22	3	US-09-112-206-38	Sequence 38, Appli	
14	110	100.0	22	3	US-09-112-206-118	Sequence 118, Appli	
15	110	100.0	22	3	US-09-112-206-134	Sequence 114, Appli	
16	110	100.0	22	4	US-09-576-8224A-38	Sequence 38, Appli	
17	110	100.0	26	1	US-07-681-701-1	Sequence 1, Appli	
18	110	100.0	26	1	US-07-681-701-7	Sequence 7, Appli	
19	110	100.0	29	3	US-08-380-160-5	Sequence 5, Appli	
20	110	100.0	30	1	US-08-324-977-6	Sequence 6, Appli	
21	110	100.0	30	2	US-08-616-616-6	Sequence 6, Appli	
22	110	100.0	30	2	US-08-604-686A-6	Sequence 6, Appli	
23	110	100.0	30	3	US-09-315-850-6	Sequence 6, Appli	
24	110	100.0	30	4	US-09-497A-47	Sequence 47, Appli	
25	110	100.0	31	1	US-07-681-701-8	Sequence 8, Appli	
26	110	100.0	32	2	US-08-146-028-47	Sequence 47, Appli	
27	110	100.0	32	2	US-08-146-028-136	Sequence 136, Appli	

ALIGNMENTS

RESULT 1
US-08-466-975A-2
; Sequence 2, Application US/08466975A
; Patent No. 5910404

; GENERAL INFORMATION:
; APPLICANT: DELLEYS, ROBERT J.
; APPLICANT: POLLET, DIRK
; APPLICANT: MAETENS, GERT
; APPLICANT: VAN HEUVERSIN, HUGO

; TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
; VARIOUS INFECTIONS: ANTIBODIES TO HEPATITIS C VIRUS

; NUMBER OF SEQUENCES: 23

; CORRESPONDENCE ADDRESS:
; ADDRESSEE: NIXON & VANDERHYE P. C.
; STREET: 1100 NORTH GLEBE ROAD
; CITY: ARLINGTON
; STATE: VA
; ZIP: 22201

; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,975A
FILING DATE:
CLASSIFICATION:
PRIORITY APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP91/02409
FILING DATE: 13-DEC-1991

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/391,671
FILING DATE:
APPLICATION NUMBER: US 07/920,286
FILING DATE: 14-OCT-1992

PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 90124241.2
FILING DATE: 14-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, R.J.
REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 1407-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 7038164100
TELEFAX: 7038164100
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid

STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-08-466-975A-2

Query Match 100.0%; Score 110; DB 2; Length 20;
 Best Local Similarity 100.0%; Pred. No. 9.2e-11;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 1 PQRKTKRNTNRRPQDVKEPG 20

RESULT 2
 US-08-391-671A-2
 Sequence 2, Application US/08391671A
 Patient No. 5922532
 GENERAL INFORMATION:
 APPLICANT: DELEYNS, ROBERT J
 APPLICANT: POLLET, DIRK
 APPLICANT: MAERTENS, GEERT
 APPLICANT: VAN HEUVERWYN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 TITLE OF INVENTION: ANTIBODIES TO HEPATITIS C VIRUS
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSSEE: NIXON & VANDERHYE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201

COMPUTER READABLE FORM:
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/391,671
 FILING DATE: 21-FEB-1995
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US-07/920,286
 FILING DATE: 14-OCT-1992
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 13-DEC-1991
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 21-FEB-1995
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-OCT-1992
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 13-DEC-1991
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-OCT-1992
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 13-DEC-1991
 REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164100
 TELEFAX: 7038164100
 REFERENCE/SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-08-391-671A-2

Query Match 100.0%; Score 110; DB 2; Length 20;
 Best Local Similarity 100.0%; Pred. No. 9.2e-11;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 1 PQRKTKRNTNRRPQDVKEPG 20

RESULT 3
 US-08-467-902A-2
 Sequence 2, Application US/08467902A
 Patient No. 6007982
 GENERAL INFORMATION:
 APPLICANT: DELEYNS, ROBERT J
 APPLICANT: POLLET, DIRK
 APPLICANT: MAERTENS, GEERT
 APPLICANT: VAN HEUVERWYN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSSEE: NIXON & VANDERHYE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/467,902A
 FILING DATE:
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164100
 TELEFAX: 7038164100
 REFERENCE/SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-08-467-902A-2

Query Match 100.0%; Score 110; DB 3; Length 20;
 Best Local Similarity 100.0%; Pred. No. 9.2e-11;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 1 PQRKTKRNTNRRPQDVKEPG 20

RESULT 4
 US-09-215-265-2
 Sequence 2, Application US/09275265
 Patient No. 6287761
 GENERAL INFORMATION:
 APPLICANT: DELEYNS, ROBERT J

APPLICANT: POLLET, DIRK
 MAERTENS, GEERT
 APPLICANT: VAN HEUVERSWIN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF INVENTION: ANTIBODIES TO HEPATITIS C VIRUS
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 STREET: NIXON & VANDERHYE P.C.
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/275,265
 FILING DATE:
 CLASSIFICATION:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US/08/391,671
 FILING DATE: 21-FEB-1995
 APPLICATION NUMBER: US 07/920,286
 FILING DATE: 14-OCT-1992
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 13-DEC-1991
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164100
 TELEFAX: 7038164100
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 S-09-275-265-2
 RESULT⁵
 S-09-941-611-2
 Sequence 2, Application US/09941611
 Patent No. 6576417
 GENERAL INFORMATION:
 APPLICANT: DELEYE, ROBERT J
 POLLET, DIRK
 MAERTENS, GEERT
 VAN HEUVERSWIN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESS: NIXON & VANDERHYE P.C.,
 STREET: 1100 NORTH GLEBE ROAD,
 CITY: ALEXANDRIA,
 STATE: VA,
 COUNTRY: USA
 ZIP: 22314-3422
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/275,265
 FILING DATE:
 CLASSIFICATION:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US/08/391,671
 FILING DATE: 21-FEB-1995
 APPLICATION NUMBER: US 07/920,286
 FILING DATE: 14-OCT-1992
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 13-DEC-1991
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164100
 TELEFAX: 7038164100
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 S-09-275-265-2
 Query Match Score 110; DB 3; Length 20;
 Best Local Similarity 100.0%; Freq. No. 9.2e-11;
 Matches 20; Conservative 0; Mismatches 0; Indels 0
 Y 1 PQRKTKRNTNRPQDVKPG 20
 b 1 PQRKTKRNTNRPQDVKPG 20

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CITY: ARLINGTON
STATE: VA
COUNTRY: USA
ZIP: 22201

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent-Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/941,611
FILING DATE: 30-Aug-2001
CLASSIFICATION: Unknown>
PRIORITY DATA:
APPLICATION NUMBER: 08/391,671
FILING DATE: 1995-02-21
APPLICATION NUMBER: WO PCT/EP91/02409
FILING DATE: 13-DEC-1991
APPLICATION NUMBER: EP 90124241.2
FILING DATE: 14-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 1,87-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 7038164000
TELEFAX: 7038164100
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-941,611-2

Query Match      100 %; Score 110; DB 4; Length
Best Local Similarity 100 %; Pred. No. 9.2e-11;
Matches 20; Conservative 0; Mismatches 0; Indels 1

Qy          1 PQRKTKRNTNRQDVKPG 20
Db          1 PQRKTKRNTNRQDVKPG 20

RESULT 6
US-09-790-497A-38
Sequence 38, Application US/09790497A
Patent No. 6649735

GENERAL INFORMATION:
APPLICANT: De Leys, Robert
TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDE
TO IMMUNOLOGICALLY IMPORTANT EPTOPE
A PROCESS FOR DETERMINATION OF ANTIBIOTINYLATED PEPTIDES CORRESPONDING TO
EPTOPE, A PROCESS FOR PREPARING THE
CONTAINING THEM
FILE REFERENCE: 2752-16
CURRENT APPLICATION NUMBER: US/09/790,497A
CURRENT FILING DATE: 2001-02-23
PRIOR APPLICATION NUMBER: 09/576,824
PRIOR FILING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 08/723,425
PRIOR FILING DATE: 1996-09-30
PRIOR APPLICATION NUMBER: 09/146,028
PRIOR FILING DATE: 1993-11-22
PRIOR APPLICATION NUMBER: PCT/EP93/00517
PRIOR FILING DATE: 1993-03-08
PRIOR APPLICATION NUMBER: EP 92400598.6
PRIOR FILING DATE: 1992-03-06
NUMBER OF SEQ ID NOS: 600
SEQUENCE ID: 1
SEQUENCE LENGTH: 20
SEQUENCE:
PQRKTKRNTNRQDVKPG

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SEQ ID NO 38
LENGTH: 20
TYPE: PRT
ORGANISM: Hepatitis C virus
US-09-790-497A-38

Query Match          100.0%; Score 110; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 9.2e-11;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 PQRKTKRNTNRRPQQVKFPG 20
Db      1 PQRKTKRNTNRRPQQVKFPG 20

RESULT 7
US-08-146-028-38
Sequence 38, Application US/08146028
Patent No. 5891610
GENERAL INFORMATION:
APPLICANT: PROCESS FOR THE DETERMINATION OF PEPTIDES
TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR
TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED
TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
NUMBER OF SEQUENCES: 453

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/146,028
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: HCV
FEATURE:
NAME/KEY: Modified-site
LOCATION: 1
FEATURE:
NAME/KEY: Modified-site
LOCATION: 22
US-08-146-028-38

Query Match          100.0%; Score 110; DB 2; Length 22;
Best Local Similarity 100.0%; Pred. No. 1e-10;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 PQRKTKRNTNRRPQQVKFPG 20
Db      2 PQRKTKRNTNRRPQQVKFPG 21

RESULT 8
US-08-146-028-118
Sequence 118, Application US/08146028
Patent No. 5891640
GENERAL INFORMATION:
APPLICANT: PROCESS FOR THE DETERMINATION OF PEPTIDES
TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR
TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED
TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
NUMBER OF SEQUENCES: 453

COMPUTER READABLE FORM:

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TITLE OF INVENTION: ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...
 NUMBER OF SEQUENCES: 453
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHUYE, P.C.
 STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
 CITY: Arlington
 STATE: VA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/723,425A
 FILING DATE:
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-13
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 703-816-4000
 TELEFAX: 703-816-4100
 INFORMATION FOR SEQ ID NO: 118:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 22 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: Peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Xaa is absent
 LOCATION: 1
 FEATURE:
 NAME/KEY: Xaa is absent
 LOCATION: 22
 US-08-723-425A-118

Query Match Score 110; DB 3; Length 22;
 Best Local Similarity 100.0%; Pred. No. 1e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qty 1 PQRKTKRNTNRPQDVKPG 20
 Db 2 PQRKTKRNTNRPQDVKPG 21

RESULT 12
 US-08-723-425A-134
 Sequence 134, Application US/08723425A
 Patent No. 61,65730
 GENERAL INFORMATION:
 APPLICANT: DELEY'S, ROBERT
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT PEPTIDES AND THEIR USE IN A PROCESS FOR DETERMINATION OF PEPTIDES CORRESPONDING ...
 TITLE OF INVENTION: ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...
 NUMBER OF SEQUENCES: 453
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHUYE, P.C.
 STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
 CITY: Arlington
 STATE: VA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS

RESULT 11
 US-08-723-425A-118
 Sequence 118, Application US/08723425A
 Patent No. 61,65730
 GENERAL INFORMATION:
 APPLICANT: DELEY'S, ROBERT
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT PEPTIDES AND THEIR USE IN A PROCESS FOR DETERMINATION OF PEPTIDES CORRESPONDING ...
 TITLE OF INVENTION: ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...
 NUMBER OF SEQUENCES: 453
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHUYE, P.C.
 STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
 CITY: Arlington
 STATE: VA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS

RESULT 10
 US-08-723-425A-38
 Sequence 38, Application US/08723425A
 GENERAL INFORMATION:
 APPLICANT: DELEY'S, ROBERT
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT PEPTIDES AND THEIR USE IN A PROCESS FOR DETERMINATION OF PEPTIDES CORRESPONDING ...
 TITLE OF INVENTION: ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...
 NUMBER OF SEQUENCES: 453
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHUYE, P.C.
 STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
 CITY: Arlington
 STATE: VA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/723,425A
 FILING DATE:
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-13
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 703-816-4000
 TELEFAX: 703-816-4100
 INFORMATION FOR SEQ ID NO: 38:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 22 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Modified-site
 LOCATION: 1
 FEATURE:
 NAME/KEY: Modified-site
 LOCATION: 22
 US-08-723-425A-38

Query Match Score 110; DB 3; Length 22;
 Best Local Similarity 100.0%; Pred. No. 1e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qty 1 PQRKTKRNTNRPQDVKPG 20
 Db 2 PQRKTKRNTNRPQDVKPG 21

SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/723,425A

FILING DATE:

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: SADOFF, B.J.

REGISTRATION NUMBER: 36,663

REFERENCE/DOCKET NUMBER: 1487-13

TELECOMMUNICATION INFORMATION:

TELEPHONE: 703-816-4000

TELEFAX: 703-816-4100

INFORMATION FOR SEQ ID NO: 134:

SEQUENCE CHARACTERISTICS:

LENGTH: 22 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ORIGINAL SOURCE:

INDIVIDUAL ISOLATE: HCV

FEATURE:

NAME/KEY: Xaa is absent

LOCATION: 1

FEATURE:

NAME/KEY: Xaa is absent

LOCATION: 22

US-08-723,425A-134

Query Match 100.0%; Score 110; DB 3; Length 22;
 Best Local Similarity 100.0%; Pred. No. 1e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20

Db 2 PQRKTKRNTNRRPQDVKFPG 21

RESULT 13

US-09-112-206-38

Sequence 38, Application US/09112206

Patent No. 6210903

GENERAL INFORMATION:

APPLICANT:

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES

TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR

TITLE OF INVENTION: BIOTINYLATED

TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED

TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPEs,

TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM

NUMBER OF SEQUENCES: 453

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

OPERATING SYSTEM: IBM PC Compatible

SOFTWARE: PC-DOS/MS-DOS

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/112,206

FILING DATE:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/146,028

FILING DATE:

INFORMATION FOR SEQ ID NO: 38:

SEQUENCE CHARACTERISTICS:

LENGTH: 22 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ORIGINAL SOURCE:

INDIVIDUAL ISOLATE: HCV

FEATURE:

NAME/KEY: Xaa is absent

LOCATION: 1

FEATURE:

NAME/KEY: Xaa is absent

LOCATION: 22

US-09-112-206-118

Query Match 100.0%; Score 110; DB 3; Length 22;

Best Local Similarity 100.0%; Pred. No. 1e-10;

Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20

Db 2 PQRKTKRNTNRRPQDVKFPG 21

RESULT 15

US-09-112-206-134

Sequence 134, Application US/09112206

Patent No. 6210903

GENERAL INFORMATION:

APPLICANT:

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES

TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPEs

TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED

TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPEs,

TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
 NUMBER OF SEQUENCES: 453
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patentin Release #1.0, Version #1.25 (BPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/112,206
 PRIORITY DATE:
 APPLICATION NUMBER: US 08/146,028
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 134:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 22 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE/KEY: Xaa is absent
 LOCATION: 1
 FEATURE/KEY: Xaa is absent
 NAME/KEY: Xaa is absent
 LOCATION: 22
 US-09-112-206-134

Query Match 100.0% Score 110; DB 3; Length 22;
 Best Local Similarity 100.0% Pred. No. 1e-10; Mismatches 0; Indels 0; Gaps 0;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PORKTKRNTNRRPDKVKEFG 20
 Db 2 PORKTKRNTNRRPDKVKEFG 21

RESULT 16
 US-09-576-824A-38
 Sequence 38, Application US/09576824A
 GENERAL INFORMATION:
 PATENT NO. 6667387
 APPLICANT: De Ley, Robert
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPEs AND THEIR USE IN A PROCESS FOR DETERMINATION OF ANTIBODIES OF BICHTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPEs, A PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
 FILE REFERENCE: 275-11
 CURRENT APPLICATION NUMBER: US/09/576,824A
 CURRENT FILING DATE: 2000-05-23
 PRIOR FILING DATE: 1996-09-30
 PRIOR APPLICATION NUMBER: 08/723,425
 PRIOR FILING DATE: 1993-11-22
 PRIOR APPLICATION NUMBER: PCT/EP93/00517
 PRIOR FILING DATE: 1993-03-08
 PRIOR FILING NUMBER: EP 92400598.6
 PRIOR FILING DATE: 1992-03-06
 NUMBER OF SEQ ID NOS: 600
 SOFTWARE: Patentin Ver. 2.1
 SEQ ID NO. 38
 LENGTH: 22
 ORGANISM: Hepatitis C virus
 FEATURE:
 NAME/KEY: VARIANT
 LOCATION: (1)
 OTHER INFORMATION: modified site
 NAME/KEY: VARIANT

; LOCATION: (22)
 ; OTHER INFORMATION: modified site
 ; US-09-576-824A-38

Query Match 100.0% Score 110; DB 4; Length 22;
 Best Local Similarity 100.0% Pred. No. 1e-10; Mismatches 0; Indels 0; Gaps 0;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PORKTKRNTNRRPDKVKEFG 20
 Db 2 PORKTKRNTNRRPDKVKEFG 21

RESULT 17
 US-07-681-701-1
 Sequence 1, Application US/07681701
 GENERAL INFORMATION:
 APPLICANT: Lacroix, Martial
 TITLE OF INVENTION: PEPTIDES AND MIXTURES THEREOF FOR
 NUMBER OF INVENTION:
 NUMBER OF SEQUENCES: 17

Query Match 100.0% Score 110; DB 1; Length 26;
 Best Local Similarity 100.0% Pred. No. 1.2e-10; Mismatches 0; Indels 0; Gaps 0;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PORKTKRNTNRRPDKVKEFG 20
 Db 7 PORKTKRNTNRRPDKVKEFG 26

RESULT 18
 US-07-681-701-7
 Sequence 7, Application US/07681701
 GENERAL INFORMATION:
 APPLICANT: Lacroix, Martial
 TITLE OF INVENTION: PEPTIDES AND MIXTURES THEREOF FOR
 NUMBER OF INVENTION:
 NUMBER OF SEQUENCES: 17

CORRESPONDENCE ADDRESS:
 STREET: FISH & NEAVE
 CITY: New York
 STATE: New York
 COUNTRY: USA
 ZIP: 10022
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/07/681,701
 FILING DATE: 19910405
 CLASSIFICATION: 530
 ATTORNEY/AGENT INFORMATION:
 NAME: Haley Jr., James F.
 REGISTRATION NUMBER: 27,794
 REFERENCE/DOCKET NUMBER: IAF-10
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (212) 715-0742
 TELEFAX: (212) 715-0673
 TELEX: 14-8367
 INFORMATION FOR SEQ ID NO: 7:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 26 amino acids
 TYPE: AMINO ACID
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-07-681-701-7

Query Match 100.0%; Score 110; DB 1; Length 26;
 Best Local Similarity 100.0%; Pred. No. 1.2e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 2 PQRKTKRNTNRRPQDVKFPG 21

RESULT 19
 US-08-380-160-5
 Sequence 5, Application US/08380160
 Patent No. 6235284
 GENERAL INFORMATION:
 APPLICANT: DALBON, Pascal
 APPLICANT: JOLIVET, Michel
 TITLE OF INVENTION: SYNTHETIC POLYPEPTIDES BELONGING TO THE HEPATITIS C VIRUS (HCV) AND WHICH CAN BE USED ESPECIALLY NUMBER OF SEQUENCES: 12
 TITLE OF INVENTION: FOR DETECTING THE LATTER
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: OLIFF & BERRIDGE
 STREET: P.O. Box 19928
 CITY: Alexandria
 STATE: VA
 COUNTY: USA
 ZIP: 22320
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/380,160
 FILING DATE:
 CLASSIFICATION: 530
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/057,471
 FILING DATE: 06 MAY 1993
 ATTORNEY/AGENT INFORMATION:

NAME: Berridge, William P.
 REGISTRATION NUMBER: 30,024
 REFERENCE/DOCKET NUMBER: WPB 28682
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (703) 836-6400
 TELEFAX: (703) 836-2787
 TELEX:
 INFORMATION FOR SEQ ID NO: 5:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 29 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 HYPOTHETICAL: NO
 ANTI-SENSE: NO
 FRAGMENT TYPE: N-terminal
 ORIGINAL SOURCE:
 ORGANISM: Human Hepatitis C Virus
 US-08-380-160-5

Query Match 100.0%; Score 110; DB 3; Length 29;
 Best Local Similarity 100.0%; Pred. No. 1.4e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 1 PQRKTKRNTNRRPQDVKFPG 20

RESULT 20
 US-08-34-97-6
 Sequence 6, Application US/08324977
 Patent No. 5747339
 GENERAL INFORMATION:
 APPLICANT: OKAYAMA, Hiroto
 APPLICANT: FUKE, Isao
 APPLICANT: MORI, Chiaki
 APPLICANT: TAKAMIZAWA, Akahisa
 APPLICANT: YOSHIDA, Iwao
 TITLE OF INVENTION: NON-A, NON-B HEPATITIS VIRUS GENOMIC
 NUMBER OF SEQUENCES: 50
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Armstrong, Westerman, Hattori, McElland &
 STREET: Naughton
 CITY: Washington
 STATE: D.C.
 COUNTRY: U.S.A.
 ZIP: 20006
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Diskette, 3.5 in, 1.44Mb
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS, Version 5.0
 SOFTWARE: ASCII
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/324,977
 FILING DATE: 18-OCT-1994
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 2-167466
 FILING DATE: 25-JUN-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 2-230921
 FILING DATE: 31-AUG-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 2-305605
 FILING DATE: 09-NOV-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/095,706
 FILING DATE: 30-JUL-1993
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 07/769,996

FILING DATE: 02-OCT-1991
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 07/635,451
 FILING DATE: 28-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: Stevens-Smith, Theresa M.
 REGISTRATION NUMBER: 36,281
 REFERENCE/DOCKET NUMBER: 900703D
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (202) 659-2930
 TELEX: 440142
 FAX: (202) 887-0357
 TELEFAX: (202) 887-0357
 INFORMATION FOR SEQ ID NO: 6:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 30 amino acids
 TOPOLogy: linear
 MOLECULE TYPE: protein
 US-08-324-977-6

Query Match 100.0%; Score 110; DB 1; Length 30;
 Best Local Similarity 100.0%; Pred. No. 1.e-10;
 Matches 20; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 21
 US-08-384-616-6
 Sequence 6, Application US/08384616
 Patent No. 5847101

GENERAL INFORMATION:
 APPLICANT: OKAYAMA, Hiroto
 APPLICANT: FUKE, Isao
 APPLICANT: MORI, Chisato
 APPLICANT: TAKAMIZAWA, Akahisa
 APPLICANT: YOSHIDA, Iwao
 TITLE OF INVENTION: NON-A, NON-B HEPATITIS VIRUS GENOMIC
 NUMBER OF SEQUENCES: 50
 ADDRESS: Armstrong, Westerman, Hattori, McLeland &
 ADDRESS: Naughton, D.C.
 STREET: 1725 K St. N.W. Suite 1000
 CITY: Washington
 STATE: U.S.A.
 ZIP: 20006

COMPUTER READABLE FORM:
 MEDIUM TYPE: Diskette, 3.5 in, 1.44Mb
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS, Version 5.0
 SOFTWARE: ASCII

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/384,616
 FILING DATE:
 CLASSIFICATION: 424
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 07/769,996
 FILING DATE: 02-OCT-1991
 APPLICATION NUMBER: JP 2-167466
 FILING DATE: 25-JUN-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 2-230921
 FILING DATE: 31-AUG-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 2-305605
 FILING DATE: 09-NOV-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 07/769,996
 FILING DATE: 02-OCT-1991
 PRIOR APPLICATION DATA:

ATTORNEY/AGENT INFORMATION:
 NAME: Stevens-Smith, Theresa M.
 REGISTRATION NUMBER: 36,281
 REFERENCE/DOCKET NUMBER: 900703B
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (202) 659-2930
 TELEX: 440142
 INFORMATION FOR SEQ ID NO: 6:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 30 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-08-384-616-6

Query Match 100.0%; Score 110; DB 2; Length 30;
 Best Local Similarity 100.0%; Pred. No. 1.e-10;
 Matches 20; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 22
 US-08-904-686A-6
 Sequence 6, Application US/08904686A
 Patent No. 5998130

GENERAL INFORMATION:
 APPLICANT: OKAYAMA, Hiroto
 APPLICANT: FUKE, Isao
 APPLICANT: MORI, Chisato
 APPLICANT: TAKAMIZAWA, Akahisa
 APPLICANT: YOSHIDA, Iwao
 TITLE OF INVENTION: NON-A, NON-B HEPATITIS VIRUS GENOMIC
 NUMBER OF SEQUENCES: 50
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Armstrong, Westerman, Hattori, McLeland &
 ADDRESS: Naughton, D.C.
 STREET: 1725 K St. N.W. Suite 1000
 CITY: Washington
 STATE: D.C.
 ZIP: 20006

COMPUTER READABLE FORM:
 MEDIUM TYPE: Diskette, 3.5 in, 1.44Mb
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS, Version 5.0
 SOFTWARE: ASCII

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/904,686A
 FILING DATE: 25-JUN-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 2-230921
 FILING DATE: 31-AUG-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/324,977
 FILING DATE: 18-OCT-1994
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 2-167466
 FILING DATE: 01-AUG-1997
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/324,977
 FILING DATE: 30-JUL-1993
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 07/769,996
 FILING DATE: 09-NOV-1990
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/099,706
 FILING DATE: 30-JUL-1993
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 07/769,996
 FILING DATE: 02-OCT-1991
 PRIOR APPLICATION DATA:

```

APPLICATION NUMBER: US 07/635,451
FILING DATE: 28-DEC-1990
ATTORNEY/AGENT INFORMATION:
    NAME: McLeland, Le-Nhung
    REGISTRATION NUMBER: 31,541
    REFERENCE/DOCKET NUMBER: 900703G
TELECOMMUNICATION INFORMATION:
    TELEPHONE: (202) 659-2330
    TELEFAX: (202) 887-0357
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-904-686A-6

Query Match          100.0%; Score 1
Best Local Similarity 100.0%; Pred. N
Matches   20; Conservative 0; Mismatch
Qy          1 PQRKTKRNTNRPQDVKPG 20
Db          7 PQRKTKRNTNRPQDVKPG 26

```

RESULT 23

US-09-315-850-6	Sequence 6, Application US/09315850
	Patent No. 6217872
GENERAL INFORMATION:	
APPLICANT:	OKAYAMA, Hiroto
APPLICANT:	FUKE, Isao
APPLICANT:	MORI, Chisato
APPLICANT:	TAKIMIZAWA, Akahisa
APPLICANT:	YOSHIDA, Iwao
TITLE OF INVENTION: NON-A, NON-B HEPATITIS	
TITLE OF INVENTION: CDNA AND ANTIGEN	
NUMBER OF SEQUENCES:	50
CORRESPONDENCE ADDRESS:	
ADDRESSEE:	Armstrong, Westerman, H
STREET:	1725 K ST. N.W. Suite 10000
CITY:	Washington
STATE:	D.C.
COUNTRY:	U.S.A.
ZIP:	20006
COMPUTER READABLE FORM:	
MEDIUM TYPE: Diskette, 3.5 in, 1.4	
COMPUTER: IBM PC compatible	
OPERATING SYSTEM: PC-DOS/MS-DOS, V	
SOFTWARE: ASCII	
CURRENT APPLICATION DATA:	
APPLICATION NUMBER:	US/09/315,850
FILING DATE:	
PRIOR APPLICATION DATA:	
APPLICATION NUMBER:	US/08/904,686
FILING DATE:	01-AUG-1997
APPLICATION NUMBER:	US 08/324,977
FILING DATE:	18-OCT-1994
PRIOR APPLICATION DATA:	
APPLICATION NUMBER:	JP 2-167466
FILING DATE:	25-JUN-1990
PRIOR APPLICATION DATA:	
APPLICATION NUMBER:	JP 2-230921
FILING DATE:	31-AUG-1990
PRIOR APPLICATION DATA:	
APPLICATION NUMBER:	JP 2-305605
FILING DATE:	09-NOV-1990
PRIOR APPLICATION DATA:	
APPLICATION NUMBER:	US 08/099,705
FILING DATE:	30-JUL-1993
PRIOR APPLICATION DATA:	

```

; APPLICATION NUMBER: US 07/769,996
; FILING DATE: 02-OCT-1991
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: US 07/635,451
;     FILING DATE: 28-DEC-1990
;     ATTORNEY/AGENT INFORMATION:
;       NAME: McLeland, Le-Nhung
;       REGISTRATION NUMBER: 31,541
;       REFERENCE/DOCKET NUMBER: 900703G
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE: (202) 659-2930
;       TELEFAX: (202) 887-0357
;     INFORMATION FOR SEQ ID NO: 6:
;       SEQUENCE CHARACTERISTICS:
;         LENGTH: 30 amino acids
;         TYPE: amino acid
;         TOPOLOGY: linear
;       MOLECULE TYPE: protein
; US-09-315-850-6

Query Match          100.0%; Score 1
Best Local Similarity 100.0%; Pred. N
Matches 20; Conservative 0; Mismatches 0

Oy      1 PQRKTRKRNRRPQDVKEPG 20
Db      7 PQRKTRKRNRRPQDVKEPG 26

RESULT 24
US-09-750-497A-47
Sequence 47 Application US/09790497A
; Patent No. 649735

; GENERAL INFORMATION:
;   APPLICANT: De Leys, Robert
;   TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF IMMUNOLOGICALLY ACTIVE PROTEINS
;   DESCRIPTION: A PROCESS FOR DETERMINING THE IMMUNOLOGICAL ACTIVITY OF PROTEINS, ESPECIALLY BIOTINYLATED PEPTIDES, BY DETERMINING THE EPITOPE(S), A PROCESS FOR THE DETERMINATION OF THE AMINO ACID SEQUENCES CONTAINING THEM
;   FILE REFERENCE: 2752.16
;   CURRENT APPLICATION NUMBER: US/09/790,4
;   CURRENT FILING DATE: 2001-02-23
;   PRIOR APPLICATION NUMBER: 09/576,824
;   PRIOR FILING DATE: 2000-05-23
;   PRIOR APPLICATION NUMBER: 08/723,425
;   PRIOR FILING DATE: 1996-09-17
;   PRIOR APPLICATION NUMBER: 09/146,028
;   PRIOR FILING DATE: 1993-11-22
;   PRIOR APPLICATION NUMBER: PCT/EP93/0051
;   PRIOR FILING DATE: 1993-03-08
;   PRIOR APPLICATION NUMBER: EP 92400598.6
;   PRIOR FILING DATE: 1992-03-06
;   NUMBER OF SEQ ID NOS: 600
;   SOFTWARE: PatentIn Ver. 2.1
;   SEQ ID NO: 47
;   LENGTH: 30
;   TYPE: PRT
;   ORGANISM: Hepatitis C virus
; US-09-750-497A-47

Query Match          100.0%; Score 1
Best Local Similarity 100.0%; Pred. N
Matches 20; Conservative 0; Mismatches 0

Oy      1 PQRKTRKRNRRPQDVKEPG 20
Db      5 PQRKTRKRNRRPQDVKEPG 24

RESULT 25
US-07-681-701-8

```

Sequence 8, Application US/07681701
 Patent No. 5574132

GENERAL INFORMATION:

APPLICANT: Lacroix, Martial
TITLE OF INVENTION: PEPTIDES AND MIXTURES THEREOF FOR CORRESPONDENCE ADDRESS: NUMBER OF SEQUENCES: 17

ADDRESSEE: FISH & NEAVE STREET: 875 Third Avenue CITY: New York STATE: New York COUNTRY: USA ZIP: 10022

COMPUTER READABLE FORM:

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/681,701
FILING DATE: 19910405
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:

NAME: Haley Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: IAF-10
TELECOMMUNICATION INFORMATION:

TELEPHONE: (212) 715-0742
TELEX: 14-8367
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 31 amino acids
TYPE: AMINO ACID
STRANDEDNESS: Single
TOPOLOGY: linear
MOLECULE TYPE: peptide

IS-07-681-701-8

RESULT 26
IS-08-146-028-47
Sequence 47, Application US/08146028
Patent No. 5891640

GENERAL INFORMATION:

APPLICANT: PROCESS FOR THE DETERMINATION OF MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25 (EPIC)
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/146, 028
INFORMATION FOR SEQ ID NO: 47:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
TOPOLOGY: linear

```

; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; INDIVIDUAL ISOLATE: HCV
; FEATURE:
; NAME/KEY: Modified-site
; LOCATION: 1
; US-08-146-028-47

Query Match Similarity 100.0%; Score 110; DB 2; Length 32;
Best Local Similarity 100.0%; Pred. No. 1.5e-10;
Matches 20; Conservative 0; Mismatches 0; Indels 0

Qy      1 PQRKTKRNTNRPQDVKEFG 20
Db      6 PQRKTKRNTNRPQDVKEFG 25

RESULT 27
US-08-146-028-136
; Sequence 136, Application US/08146028
; Patent No. 5891640
; GENERAL INFORMATION:
; APPLICANT: PROCESS FOR THE DETERMINATION OF PEPTIDES
; CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
; IN A PROCESS FOR DETERMINATION OF ANTIBODIES
; PEPTIDES CORRESPONDING TO IMMUNOLOGICAL COMPOSITIONS
; TITLE OF INVENTION:
; NUMBER OF SEQUENCES: 453
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/146,028
; ID: SEQ ID NO: 136:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; INDIVIDUAL ISOLATE: HCV
; FEATURE:
; NAME/KEY: Xaa is absent
; LOCATION: 1
; FEATURE:
; NAME/KEY: Xaa is absent
; LOCATION: 32
; US-08-146-028-136

Query Match Similarity 100.0%; Score 110; DB 2; Length 32;
Best Local Similarity 100.0%; Pred. No. 1.5e-10;
Matches 20; Conservative 0; Mismatches 0; Indels 0

Qy      1 PQRKTKRNTNRPQDVKEFG 20
Db      6 PQRKTKRNTNRPQDVKEFG 25

RESULT 28...
US-08-723-425A-47
; Sequence 47, Application US/08723425A
; Patent No. 6165730
; GENERAL INFORMATION:
; APPLICANT: DELETS, ROBERT
; TITLE OF INVENTION:
; PROCESS FOR THE DETERMINATION OF

```

TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR USE IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...
 TITLE OF INVENTION: 453
 NUMBER OF SEQUENCES: 453
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE, P.C.
 STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
 CITY: Arlington
 STATE: Va
 COUNTRY: USA
 ZIP: 22201

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 TELECOMMUNICATION INFORMATION:
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/723,425A
 FILING DATE:
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-13
 TELEPHONE: 703-816-4000
 TELEXFAX: 703-816-4100
 INFORMATION FOR SEQ ID NO: 136:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 32 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: Peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Xaa is absent
 LOCATION: 1
 FEATURE:
 NAME/KEY: Xaa is absent
 LOCATION: 32

Query Match 100.0%; Score 110; DB 3; Length 32;
 Best Local Similarity 100.0%; Pred. No. 1.5e-10; Indels 0; Gaps 0;
 Matches 20; Conservative 0; Mismatches 0; Pat. No. 6210903
 ; Sequence 47, Application US/09112206
 ; Patent No. 6210903
 ; GENERAL INFORMATION:
 ; APPLICANT:
 ; TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES
 ; CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR
 ; USE IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED
 ; PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE,
 ; PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
 ; NUMBER OF SEQUENCES: 453
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: PatentIn Release #1.0, Version #1.25 (BPO)

RESULT 29
 US-08-723-425A-136
 ; Sequence 136, Application US/08723425A
 ; Patent No. 6165730

GENERAL INFORMATION:
 APPLICANT: DELEY, ROBERT
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR USE IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...
 NUMBER OF SEQUENCES: 453
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE, P.C.
 STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
 CITY: Arlington
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk

NAME/KEY: Modified-site
 LOCATION: 1
 FEATURE: Modified-site
 NAME/KEY: Modified-site
 LOCATION: 32
 US-09-112-206-47

Query Match Best Local Similarity 100.0%; Score 110; DB 3; Length 32;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKPG 20
 Db 6 PQRKTKRNTNRRPQDVKPG 25

RESULT 31
 US-09-112-206-136
 ; Sequence 136, Application US/09112206
 ; Patent No. 6210503
 ; GENERAL INFORMATION:
 ; APPLICANT: PROCESS FOR THE DETERMINATION OF PEPTIDES
 ; TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR
 ; TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED
 ; TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE,
 ; TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
 ; NUMBER OF SEQUENCES: 453
 ; COMPUTER READABLE FORM:
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: PatentIn Release #1.0, version #1.25 (BPO)

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/112,206
 FILING DATE:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/146,028
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 136:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 32 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Xaa is absent
 LOCATION: 1
 ;
 NAME/KEY: Xaa is absent
 LOCATION: 32
 US-09-112-206-136

Query Match Best Local Similarity 100.0%; Score 110; DB 3; Length 32;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKPG 20
 Db 6 PQRKTKRNTNRRPQDVKPG 25

RESULT 32
 US-09-790-497A-136
 ; Sequence 136, Application US/09790497A
 ; Patent No. 6649735
 ; GENERAL INFORMATION:
 ; APPLICANT: De Leyb, Robert
 ; TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
 ; TITLE OF INVENTION: TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR USE IN
 ; TITLE OF INVENTION: A PROCESS FOR DETERMINATION OF ANTIBODIES OF
 ; TITLE OF INVENTION: BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
 ; TITLE OF INVENTION: EPITOPE, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
 ; TITLE OF INVENTION: CONTAINING THEM
 ; FILE REFERENCE: 2752-16
 ; CURRENT APPLICATION NUMBER: US/09/790,497A
 ; CURRENT FILING DATE: 2001-02-23
 ; PRIOR APPLICATION NUMBER: 09/576,824
 ; PRIOR FILING DATE: 1996-09-30
 ; PRIOR APPLICATION NUMBER: 09/146,028
 ; PRIOR FILING DATE: 1993-11-22
 ; PRIOR APPLICATION NUMBER: PCT/EP93/00517
 ; PRIOR FILING DATE: 1993-03-08
 ; PRIOR APPLICATION NUMBER: EP 92400598-6
 ; PRIOR FILING DATE: 1992-03-06
 ; NUMBER OF SEQ ID NOS: 600
 ; SEQ ID NO 136
 ; LENGTH: 32
 ; TYPE: PRT
 ; ORGANISM: Hepatitis C virus
 US-09-790-497A-136

Query Match Best Local Similarity 100.0%; Score 110; DB 4; Length 32;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKPG 20

Db 7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 34 ; NUMBER OF SEQ ID NOS: 600

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO: 136

; LENGTH: 32

; TYPE: PRT

; ORGANISM: Hepatitis C virus

US-09-576-824A-136

GENERAL INFORMATION:

; APPLICANT: De Leyva, Robert

; TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING

; TO IMMUNOLOGICALLY IMPORTANT EPITOPEs AND THEIR USE IN

; A PROCESS FOR DETERMINATION OF ANTIBODIES OF

; BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT

; EPITOPEs, A PROCESS FOR PREPARING THEM AND COMPOSITIONS

; TITLE OF INVENTION: CONTAINING THEM

FILE REFERENCE: 2752-11

CURRENT APPLICATION NUMBER: US/09/576,824A

CURRENT FILING DATE: 2000-05-23

PRIOR APPLICATION NUMBER: 08/723,425

PRIOR FILING DATE: 1996-09-30

PRIOR APPLICATION NUMBER: 09/146,028

PRIOR FILING DATE: 1993-11-22

PRIOR APPLICATION NUMBER: PCT/EP93/00517

PRIOR FILING DATE: 1993-03-08

PRIOR APPLICATION NUMBER: EP 92400598.6

PRIOR FILING DATE: 1992-03-06

NUMBER OF SEQ ID NOS: 600

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO: 47

LENGTH: 32

TYPE: PRT

ORGANISM: Hepatitis C virus

FEATURE:

NAME/KEY: VARIANT

LOCATION: (1)

OTHER INFORMATION: modified site

NAME/KEY: VARIANT

LOCATION: (32)

OTHER INFORMATION: modified site

US-09-576-824A-47

Query Match 100.0%; Score 110; DB 4; Length 32;

Best Local Similarity 100.0%; Pred. No. 1.5e-10; Indels 0; Gaps 0;

Db 6 PQRKTKRNTNRRPQDVKFPG 25

Query Match 100.0%; Score 110; DB 4; Length 32;

Best Local Similarity 100.0%;保守型 0; Mismatches 0; Indels 0; Gaps 0;

Db 7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 35 ; NUMBER OF SEQ ID NOS: 160

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO: 136

; LENGTH: 32

; TYPE: PRT

; ORGANISM: Human Hepatitis C Virus

US-09-576-824A-136

GENERAL INFORMATION:

; APPLICANT: De Leyva, Robert

; TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING

; TO IMMUNOLOGICALLY IMPORTANT EPITOPEs AND THEIR USE IN

; A PROCESS FOR DETERMINATION OF ANTIBODIES OF

; BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT

; EPITOPEs, A PROCESS FOR PREPARING THEM AND COMPOSITIONS

; TITLE OF INVENTION: CONTAINING THEM

FILE REFERENCE: 2752-11

CURRENT APPLICATION NUMBER: US/09/576,824A

CURRENT FILING DATE: 2000-05-23

PRIOR APPLICATION NUMBER: 08/723,425

PRIOR FILING DATE: 1996-09-30

PRIOR APPLICATION NUMBER: 09/146,028

PRIOR FILING DATE: 1993-11-22

PRIOR APPLICATION NUMBER: PCT/EP93/00517

PRIOR FILING DATE: 1992-03-08

PRIOR APPLICATION NUMBER: EP 92400598.6

PRIOR FILING DATE: 1992-03-06

Query Match 100.0%; Score 110; DB 3; Length 34;

Best Local Similarity 100.0%;保守型 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 6 PQRKTKRNTNRRPQDVKFPG 25

RESULT 37
 US-09-576-824A-402
 Sequence 402, Application US/095766824A
 Patent No. 6667878

GENERAL INFORMATION:
 APPLICANT: De Leys, Robert
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOLES AND THEIR USE IN A PROCESS FOR DETERMINATION OF ANTIBODIES OF BICONTINUED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOLES, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
 TITLE OF INVENTION: PEPTIDES CONTAINING THEM
 FILE REFERENCE: 2752-11
 CURRENT FILING DATE: 2000-05-23
 PRIOR APPLICATION NUMBER: 08/123,425
 PRIOR FILING DATE: 1996-09-30
 PRIOR APPLICATION NUMBER: 09/146,028
 PRIOR FILING DATE: 1993-11-22
 PRIOR APPLICATION NUMBER: PCT/EP93/00517
 PRIOR FILING DATE: 1993-03-08
 PRIOR APPLICATION NUMBER: EP 92400598.6
 PRIOR FILING DATE: 1992-03-06
 NUMBER OF SEQ ID NOS: 600
 SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO: 402
 LENGTH: 34
 TYPE: PRT
 ORGANISM: Hepatitis C virus

FEATURE:
 NAME/KEY: VARIANT
 LOCATION: (1)
 OTHER INFORMATION: Xaa = modified site : when present, represents an amino acid, amino group, or chemically modified other information: amino acid, amino terminus
 NAME/KEY: VARIANT
 LOCATION: (34)
 OTHER INFORMATION: Xaa = modified site : when present, represents an amino acid, OH-group, NH2-group, or a linkage other information: involving these two groups

US-09-576-824A-402

Query Match 100.0%; Score 110; DB 4; Length 34;
 Best Local Similarity 100.0%; Pred. No. 1.6e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 8 PQRKTKRNTNRRPQDVKFPG 27

RESULT 38
 US-08-380-160-8
 Sequence 8, Application US/08380160
 Patent No. 623584

GENERAL INFORMATION:
 APPLICANT: DALBON, Pascal
 APPLICANT: JOLIVET, Michel
 TITLE OF INVENTION: SYNTHETIC POLYPEPTIDES BELONGING TO THE TITLE OF INVENTION: HEPATITIS C VIRUS (HCV) AND WHICH CAN BE USED ESPECIALLY TITLE OF INVENTION: FOR DEFECTING THE LATTER
 NUMBER OF SEQUENCES: 12
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: OLIFF & BERRIDGE
 STREET: P.O. Box 19928
 CITY: Alexandria
 STATE: VA
 COUNTRY: USA

ZIP: 22320
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/380,160
 FILING DATE:

CLASSIFICATION: 530
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/057,471
 FILING DATE: 06-MAY-1993
 ATTORNEY/AGENT INFORMATION:
 NAME: Barridge, William P.
 REGISTRATION NUMBER: 30,024
 REFERENCE/DOCKET NUMBER: WPB 28682
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (703) 836-6400
 TELEX: (703) 836-2787
 INFORMATION FOR SEQ ID NO: 8:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 39 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ANTI-SENSE: NO
 FRAGMENT TYPE: N-terminal
 ORIGINAL SOURCE:
 ORGANISM: Human Hepatitis C Virus

US-08-380-160-8

Query Match 100.0%; Score 110; DB 3; Length 39;
 Best Local Similarity 100.0%; Pred. No. 1.9e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 1 PQRKTKRNTNRRPQDVKFPG 20

RESULT 39
 US-09-020-846-36

Query Match 100.0%; Score 110; DB 3; Length 39;
 Best Local Similarity 100.0%; Pred. No. 1.9e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 1 PQRKTKRNTNRRPQDVKFPG 20

RESULT 40
 US-09-020-846-36

Sequence 36, Application US/09020846
 Patent No. 6329965

GENERAL INFORMATION:
 APPLICANT: YAMAGUCHI, Keniro.
 APPLICANT: KASHIWAKUMA, Tomiko
 APPLICANT: CHIBA, Yukie
 APPLICANT: YAGI, Shintaro
 APPLICANT: HASEGAWA, Akira
 TITLE OF INVENTION: CHIMERA ANTIGEN PEPTIDE
 NUMBER OF SEQUENCES: 72
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: FOLLY & LARDNER
 STREET: 3000 K Street, N.W.
 CITY: Washington
 STATE: D.C.
 COUNTRY: U.S.A.
 ZIP: 20007-5109
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/020,846
 FILING DATE: 09-FEB-1998
 CLASSIFICATION: 424
 PRIOR APPLICATION DATA:

APPLICATION NUMBER: JP 9-027015
 FILING DATE: 10-FEB-1997
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: JP 8 - 024045
 FILING DATE: 09-FEB-1996
 ATTORNEY/AGENT INFORMATION:
 NAME: Wagner, Harold C.
 REGISTRATION NUMBER: 25,258
 REFERENCE/DOCKET NUMBER: 053466/0225
 TELECOMMUNICATION INFORMATION:
 LOCATION: (202) 672-5300
 TELEFAX: (202) 672-5399
 INFORMATION FOR SEQ ID NO: 36:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 43 amino acids
 TYPE: amino acid
 STRANDEDNESS:
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-09-020-846-36

Query Match 100.0%; Score 110; DB 4; Length 43;
 Best Local Similarity 100.0%; Pred. No. 2,1e-10;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 40
 US-08-380-160-2
 Sequence 2, Application US/08380160
 Patent No. 6235284

GENERAL INFORMATION:
 APPLICANT: DALBON, Pascal
 APPLICANT: JOLIVET, Michel
 TITLE OF INVENTION: SYNTHETIC POLYPEPTIDES BELONGING TO THE TITLUS OF INVENTION: HEPATITIS C VIRUS (HCV) AND WHICH CAN BE USED ESPECIALLY NUMBER OF SEQUENCES: 12
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: OLIVE & BERRIDGE
 STREET: P.O. Box 19328
 CITY: Alexandria
 STATE: VA
 COUNTRY: USA
 ZIP: 22320

COMPUTER READABLE FORM:
 COMPUTER TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent In Release #1.0, Version #1.1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/380,160
 FILING DATE:
 CLASSIFICATION: 530
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/057,471
 FILING DATE: 06-MAY-1993
 ATTORNEY/AGENT INFORMATION:
 NAME: Berridge, William P.
 REGISTRATION NUMBER: 30,124
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (703) 836-6400
 TELEX: (703) 336-2787

INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 44 amino acids
 TYPE: amino acid
 STRANDEDNESS: single

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OM protein - protein search, using bw model.

Run on: July 20, 2004, 09:34:06 ; Search time 40 Seconds
(without alignments)

Scoring table: BIOSUM62
Gapext 10.0 , Gapext 0.5

Searched: 1285356 seqs, 312560742 residues

Total number of hits satisfying chosen parameters: 1285356

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications-AA+
1: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US07_PUBCOMB.pep:
2: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US06_NEW_PUB.pep:
3: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US06_NEW_PUB.pep:
4: /cn2_6_ptodata/1/pubbaa/1/pubbaa/PCT_NEW_PUB.pep:
5: /cn2_6_ptodata/1/pubbaa/1/pubbaa/PCTUS_PUBCOMB.pep:
6: /cn2_6_ptodata/1/pubbaa/1/pubbaa/PCTUS_PUBCOMB.pep:
7: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US08_NEW_PUB.pep:
8: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US08_PUBCOMB.pep:
9: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US09_PUBCOMB.pep:
10: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US09B_PUBCOMB.pep:
11: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US09C_PUBCOMB.pep:
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13: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US10A_PUBCOMB.pep:
14: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US10B_PUBCOMB.pep:
15: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US10C_PUBCOMB.pep:
16: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US60_NEW_PUB.pep:
17: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US60_NEW_PUB.pep:
18: /cn2_6_ptodata/1/pubbaa/1/pubbaa/US60_PUBCOMB.pep:
Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	Match	Length	DB ID	Description
1	106	100.0	20	9	US-09-941-611-1	Sequence 1, Appli
2	106	100.0	20	14	US-10-044-995-1	Sequence 1, Appli
3	106	100.0	450	12	US-10-165-165-181	Sequence 181, Appli
4	106	100.0	2894	9	US-09-941-611-23	Sequence 23, Appli
5	106	100.0	2894	14	US-10-044-995-23	Sequence 23, Appli
6	104	98.1	100	12	US-10-165-165-32	Sequence 232, Appli
7	104	98.1	137	9	US-09-851-138-46	Sequence 46, Appli
8	99	93.4	30	12	US-10-295-734-408	Sequence 408, Appli
9	99	93.4	48	9	US-09-851-138-22	Sequence 22, Appli
10	99	93.4	53	12	US-10-431-587-3	Sequence 3, Appli
11	99	93.4	53	12	US-10-431-587-10	Sequence 10, Appli
12	99	93.4	53	12	US-10-431-587-11	Sequence 11, Appli
13	99	93.4	53	12	US-10-431-587-12	Sequence 12, Appli
14	99	93.4	73	12	US-10-431-587-13	Sequence 13, Appli
15	99	93.4	74	9	US-09-851-138-10	Sequence 10, Appli

ALIGNMENTS

RESULT 1
US-09-941-611-1
; Sequence 1, Application US/09941611
; Patent No. US200210664041
; GENERAL INFORMATION:
; APPLICANT: DELEYE, ROBERT J
; POLLET, DIRK
; MAERTENS, GERT
; VAN HEUVERSURN, HUGO
; TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: NIXON & VANDERHYE P.C.
; STREET: 1100 NORTH GLEBE ROAD
; CITY: ARLINGTON
; STATE: VA
; COUNTRY: USA
; ZIP: 22201
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.3.0

CURRENT APPLICATION DATA:
APPLYING DATE: 08/09/1991
APPLYING NUMBER: US/09/941-611
FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
PRIORITY DATA:
APPLICATION NUMBER: 08/391,671
FILING DATE: 1995-02-21
APPLYING NUMBER: WO PCT/EP91/02409
FILING DATE: 13-DEC-1991
APPLYING NUMBER: EP 90124241.2
FILING DATE: 14-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,663

REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 TELEFAX: 7038164100
 INFORMATION FOR SEQ ID NO: 1:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: Linear
 MOLECULE TYPE: Peptide
 SEQUENCE DESCRIPTION: SEQ ID NO: 1:
 US-09-941-611-1

Query Match 100.0%; Score 106; DB 9; Length 20;
 Best Local Similarity 100.0%; Pred. No. 4.1e-08;
 Matches 20; Conservative 0; N mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPQRTKTRNTNRRPQ 20
 Db 1 MSTIPKPQRTKTRNTNRRPQ 20

RESULT 2
 US-10-044-995-1
 Sequence 1, Application US/10044995
 Publication No. US20010049685A1
 GENERAL INFORMATION:
 APPLICANT: DELEYS, ROBERT J
 POLLET, DIRK
 MAERTENS, GEERT
 VAN HEUVERSWN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.3.0
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/10/044,995
 FILING DATE: 15-Jan-2002
 CLASSIFICATION: <Unknown>
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/391,671
 FILING DATE: <Unknown>
 APPLICATION NUMBER: US 07/920,286
 FILING DATE: 14-OCT-1992
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 13-DEC-1991
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 INFORMATION FOR SEQ ID NO: 1:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: Linear

MOLECULE TYPE: peptide
 SEQUENCE DESCRIPTION: SEQ ID NO: 1:
 US-10-044-995-1
 Query Match 100.0%; Score 106; DB 14; Length 20;
 Best Local Similarity 100.0%; Pred. No. 4.1e-08;
 Matches 20; Conservative 0; N mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPQRTKTRNTNRRPQ 20
 Db 1 MSTIPKPQRTKTRNTNRRPQ 20

RESULT 3
 US-10-651-165-181
 Sequence 18, Application US/10651165
 Publication No. US20040047877A1
 GENERAL INFORMATION:
 APPLICANT: LEROUX-ROELS, Geert
 APPLICANT: DELLEYS, Robert
 APPLICANT: MAERTENS, Geert
 TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C
 FILE REFERENCE: 2551-94
 CURRENT APPLICATION NUMBER: US/10/651,165
 CURRENT FILING DATE: 2003-09-02
 PRIORITY APPLICATION NUMBER: US/08/974,690C
 PRIORITY FILING DATE: 1997-11-19
 PRIORITY APPLICATION NUMBER: PCT/EP94/03555
 PRIORITY FILING DATE: 1994-10-28
 PRIORITY APPLICATION NUMBER: EP 93402718.6
 PRIORITY FILING DATE: 1993-11-04
 NUMBER OF SEQ ID NO: 286
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 181
 LENGTH: 450
 TYPE: PRT
 ORGANISM: hepatitis C virus
 US-10-651-165-181

Query Match 100.0%; Score 106; DB 12; Length 450;
 Best Local Similarity 100.0%; Pred. No. 8.7e-07;
 Matches 20; Conservative 0; N mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPQRTKTRNTNRRPQ 20
 Db 1 MSTIPKPQRTKTRNTNRRPQ 20

RESULT 4
 US-09-941-611-23
 Sequence 23, Application US/09541611
 Patient No. US2002106640A1
 GENERAL INFORMATION:
 APPLICANT: DELEYS, ROBERT J
 POLLET, DIRK
 MAERTENS, GEERT
 VAN HEUVERSWN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.3.0
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/10/044,995
 FILING DATE: 15-Jan-2002
 CLASSIFICATION: <Unknown>

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/941,611
 FILING DATE: 30-Aug-2001
 CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/391,671
 FILING DATE: 1995-02-21
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 13-DEC-1991
 APPLICATION NUMBER: EP 9012441.2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 TELEX/FAX: 7038164100
 INFORMATION FOR SEQ ID NO: 23:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-5
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2894 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: Peptide
 HYPOTHETICAL: NO
 ANTI-SENSE: NO
 SEQUENCE DESCRIPTION: SEQ ID NO: 23:
 US-10-044-995-23

Query Match Score 106; DB 9; Length 2894;
 Best Local Similarity 100.0%; Pred. No. 5.5e-06;
 Matches 20; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;

RESULT 5
 US-10-044-995-23
 Sequence 23, Application US/10044995
 Publication No. US20030049685AA.
 GENERAL INFORMATION:
 APPLICANT: DELEYNS, ROBERT J
 POLLER, DIRK
 MARTENS, GEERT
 VAN HEUVERSINN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/10/044,995
 FILING DATE: 15-Jan-2002
 CLASSIFICATION: <Unknown>
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/391,671
 FILING DATE: <Unknown>
 APPLICATION NUMBER: US 07/920,286
 FILING DATE: 14-OCT-1992
 APPLICATION NUMBER: WO PCT/EP91/02409

FILING DATE: 13-DEC-1991
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 TELEX/FAX: 7038164100
 INFORMATION FOR SEQ ID NO: 23:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 2894 amino acids
 TYPE: amino acid
 STRANDEDNESS: Single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ANTI-SENSE: NO
 SEQUENCE DESCRIPTION: SEQ ID NO: 23:
 US-10-044-995-23

Query Match Score 106; DB 14;
 Best Local Similarity 100.0%; Pred. No. 5.5e-06;
 Matches 20; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;

Qy 1 MSTIPKPQRTKTKNTNRRPQ 20
 Db 1 MSTIPKPQRTKTKNTNRRPQ 20

RESULT 6
 US-10-051-165-232
 Sequence 232, Application US/10651165
 Publication No. US20040047877A1
 GENERAL INFORMATION:
 APPLICANT: LEROUX-ROELS, Geert
 APPLICANT: DELEYNS, Robert
 APPLICANT: MAERTENS, Geert
 TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C
 FILE REFERENCE: 2551-94
 CURRENT APPLICATION NUMBER: US/10/651,165
 PRIORITY APPLICATION NUMBER: US/08/974,690C
 PRIORITY FILING DATE: 1997-11-19
 PRIORITY APPLICATION NUMBER: PCT/EP94/03555
 PRIORITY FILING DATE: 1994-10-28
 PRIORITY APPLICATION NUMBER: EP 93402718.6
 PRIORITY FILING DATE: 1993-11-04
 NUMBER OF SEQ ID-NOS: 286
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 232
 LENGTH: 100
 TYPE: PRT
 ORGANISM: hepatitis C virus
 US-10-051-165-232

Query Match Score 104; DB 12;
 Best Local Similarity 95.0%; Pred. No. 3.8e-07;
 Matches 19; Conservative 1; Mismatches 0;
 Indels 0; Gaps 0;

Qy 1 MSTIPKPQRTKTKNTNRRPQ 20
 Db 1 MSTIPKPQRTKTKNTNRRPQ 20

RESULT 7
 US-09-051-138-46
 Sequence 46, Application US/09851138
 Publication No. US20020183508A1
 GENERAL INFORMATION:
 APPLICANT: MAERTENS, GEERT

TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS

NUMBER OF SEQUENCES: 207

CORRESPONDENCE ADDRESS:

STREET: P. O. BOX 4433

CITY: HOUSTON

STATE: TEXAS

ZIP: 77210-4433

COMPUTER READABLE FORM:

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/851,138

FILING DATE: 09-May-2001

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/836,075

FILING DATE: <Unknown>

APPLICATION NUMBER: EP 94870166.9

FILING DATE: 21-CTC 1994

APPLICATION NUMBER: EP 95-870076.7

FILING DATE: 28-Jun 1995

ATTORNEY/AGENT INFORMATION:

NAME: KAMMERER, PATRICIA A.

REGISTRATION NUMBER: 29,775

REFERENCE/DOCKET NUMBER: INNS:004

INFORMATION FOR SEQ ID NO: 46:

SEQUENCE CHARACTERISTICS:

LENGTH: 137 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 46:

US-09-851-138-46

Query Match 98.1%; Score 104; DB 9; Length 137;

Best Local Similarity 95.0%; Pred. No. 5.2e-07;

Matches 19; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRNTRRQ 20

Db 1 MST1PKPQRTKRNTRRQ 20

RESULT 8

US-10-296-734-408

Sequence 408 Application US/10296734

Publication No. US20040054137A1

GENERAL INFORMATION:

APPLICANT: Thompson, Scott A

APPLICANT: Ramshaw, Ian A

TITLE OF INVENTION: Synthetic molecules and uses therefor

FILE REFERENCE: Savine

CURRENT APPLICATION NUMBER: US/10/296,734

PRIOR APPLICATION NUMBER: AU PQ7761/00

PRIOR FILING DATE: 2000-05-26

NUMBER OF SEQ ID NOS: 1507

SOFTWARE: PatentIn version 3.2

SEQ ID NO: 408

LENGTH: 30

TYPE: PRT

ORGANISM: Artificial

FEATURE: OTHER INFORMATION: HepC 1a segment 1

Query Match 93.4%; Score 99; DB 12; Length 30;

US-10-296-734-408

Query Match

Best Local Similarity 95.0%; Pred. No. 5.8e-07;

Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRNTRRQ 20

Db 3 MST1PKPQRTKRNTRRQ 22

RESULT 9

US-09-851-138-22

Sequence 22 Application US/09851138

Publication No. US20020183508A1

GENERAL INFORMATION:

APPLICANT: MAERTENS, GEERT

STUYVER, LIEVEN

TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS

NUMBER OF SEQUENCES: 207

CORRESPONDENCE ADDRESS:

STREET: ARNOLD, WHITE & DURKEE

CITY: HOUSTON

STATE: ARNOLD, WHITE & DURKEE

STREET: P.O. BOX 4433

CITY: HOUSTON

STATE: TEXAS

COUNTRY: USA

ZIP: 77210-4433

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/851,138

FILING DATE: 09-May-2001

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/836,075

FILING DATE: <Unknown>

APPLICATION NUMBER: EP 94870166.9

FILING DATE: 21-Oct 1994

APPLICATION NUMBER: EP 95870076.7

FILING DATE: 28-Jun 1995

ATTORNEY/AGENT INFORMATION:

NAME: KAMMERER, PATRICIA A.

REGISTRATION NUMBER: 29,775

REFERENCE/DOCKET NUMBER: INNS:004

INFORMATION FOR SEQ ID NO: 22:

SEQUENCE CHARACTERISTICS:

LENGTH: 48 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 22:

US-09-851-138-22

Query Match 93.4%; Score 99; DB 9; Length 48;

Best Local Similarity 95.0%; Pred. No. 9.3e-07;

Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRNTRRQ 20

Db 1 MST1PKPQRTKRNTRRQ 20

RESULT 10

US-10-431-587-3

Sequence 3 Application US/10431587

Publication No. US20040072267A1

GENERAL INFORMATION:

APPLICANT: BIORAD PASTEUR

TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism

FILE REFERENCE: BET 03P0456

CURRENT APPLICATION NUMBER: US/10/431,587

CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 3
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
; US-10-431-587-3

Query Match 93.4%; Score 99; DB 12; Length 53;
; Best Local Similarity 95.0%; Pred. No. 1e-06;
; Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTKNTNRRPQ 20
Db 1 MSTNPKPQRTKRTKNTNRRPQ 20

RESULT 11
US-10-431-587-10
; Sequence 10, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 10
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
; US-10-431-587-10

Query Match 93.4%; Score 99; DB 12; Length 53;
; Best Local Similarity 95.0%; Pred. No. 1e-06;
; Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTKNTNRRPQ 20
Db 1 MSTNPKPQRTKRTKNTNRRPQ 20

RESULT 12
US-10-431-587-11
; Sequence 11, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 11
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
; US-10-431-587-11

Query Match 93.4%; Score 99; DB 12; Length 53;
; Best Local Similarity 95.0%; Pred. No. 1e-06;

RESULT 13
US-10-431-587-12
; Sequence 12, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; TITLE OF INVENTION: against, an infectious microorganism
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 12
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
; US-10-431-587-12

Query Match 93.4%; Score 99; DB 12; Length 53;
; Best Local Similarity 95.0%; Pred. No. 1e-06;
; Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTKNTNRRPQ 20
Db 1 MSTNPKPQRTKRTKNTNRRPQ 20

RESULT 14
US-10-431-587-13
; Sequence 13, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; TITLE OF INVENTION: against, an infectious microorganism
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 13
; LENGTH: 73
; TYPE: PRT
; ORGANISM: Hepatitis C virus
; US-10-431-587-13

Query Match 93.4%; Score 99; DB 12; Length 73;
; Best Local Similarity 95.0%; Pred. No. 1.4e-06;
; Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTKNTNRRPQ 20
Db 1 MSTNPKPQRTKRTKNTNRRPQ 20

RESULT 15
US-09-851-138-10
; Sequence 10, Application US/098511138
; Publication No. US20020183508A1
; GENERAL INFORMATION:
; APPLICANT: MAERTENS, GBRRT

Query Match 93.4%; Score 99; DB 12; Length 53;
; Best Local Similarity 95.0%; Pred. No. 1e-06;

STUVVER, LIEVEN
TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS
NUMBER OF SEQUENCES: 207
CORRESPONDENCE ADDRESS:
ADDRESSEE: ARNOLD, WHITE & DURKEE
STREET: P.O. BOX 4433
CITY: HOUSTON
STATE: TEXAS
ZIP: 77210-4433

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/851,138
FILING DATE: 03-May-2001
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/836,075
FILING DATE: <Unknown>
APPLICATION NUMBER: EP 94870166.9
FILING DATE: 21 Oct 1994
APPLICATION NUMBER: EP 95870076.7
FILING DATE: 28 Jun 1995
ATTORNEY/AGENT INFORMATION:
NAME: KAMMERER, PATRICIA A.
REGISTRATION NUMBER: 29,775
REFERENCE/DOCKET NUMBER: INNS:004
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 74 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 10:

US-09-851-138-3

Query Match 93.4%; Score 99; DB 9; Length 74;
Best Local Similarity 95.0%; Pred. No. 1.4e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
Db 1 MSTNPKPQRKTKRNTNRRPQ 20

RESULT 16
US-10-651-165-198
Sequence 198, Application US/10651165
Publication No. US2004004787A1
GENERAL INFORMATION:
APPLICANT: LEROUX-ROELIS, Geert
APPLICANT: MAETENS, Robert
APPLICANT: MAETENS, Geert
TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPE OF HEPATITIS C VIRUS
FILE REFERENCE: 2551-94
CURRENT APPLICATION NUMBER: US/10/651,165
PRIOR FILING DATE: 1997-11-19
PRIOR APPLICATION NUMBER: PCT/EP94/03555
PRIOR FILING DATE: 1994-10-28
PRIOR APPLICATION NUMBER: EP 93402718.6
PRIOR FILING DATE: 1994-11-04
NUMBER OF SEQ ID NOS: 286
SOFTWARE: PatentIn version 3.1
SEQ ID NO: 198
LENGTH: 74
TYPE: PRT

; ORGANISM: hepatitis C virus
US-10-651-165-198

Query Match 93.4%; Score 99; DB 12; Length 74;
Best Local Similarity 95.0%; Pred. No. 1.4e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
Db 1 MSTNPKPQRKTKRNTNRRPQ 20

RESULT 17
US-10-431-587-1
Sequence 1, Application US/10431587
Publication No. US20040072267A1
GENERAL INFORMATION:
APPLICANT: BIOPAC PASTEUR
TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
FILE REFERENCE: BET 03P0456
CURRENT APPLICATION NUMBER: US/10/431,587
CURRENT FILING DATE: 2003-05-08
PRIOR APPLICATION NUMBER: FR 0205808
PRIOR FILING DATE: 2002-05-10
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 75
TYPE: PRT
ORGANISM: Hepatitis C virus
US-10-431-587-1

Query Match 93.4%; Score 99; DB 12; Length 75;
Best Local Similarity 95.0%; Pred. No. 1.e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
Db 1 MSTNPKPQRKTKRNTNRRPQ 20

RESULT 18
US-10-431-587-4
Sequence 4, Application US/10431587
Publication No. US20040072267A1
GENERAL INFORMATION:
APPLICANT: BIOPAC PASTEUR
TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
FILE REFERENCE: BET 03P0456
CURRENT APPLICATION NUMBER: US/10/431,587
CURRENT FILING DATE: 2003-05-08
PRIOR APPLICATION NUMBER: FR 0205808
PRIOR FILING DATE: 2002-05-10
NUMBER OF SEQ ID NOS: 33
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 75
TYPE: PRT
ORGANISM: Hepatitis C virus
US-10-431-587-4

Query Match 93.4%; Score 99; DB 12; Length 75;
Best Local Similarity 95.0%; Pred. No. 1.4e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
Db 1 MSTNPKPQRKTKRNTNRRPQ 20

RESULT 19

US-10-431-587-5
 ; Sequence 5, Application US/10431587
 ; Publication No. US20040072267A1
 ; GENERAL INFORMATION:
 ; APPLICANT: BIORAD PASTEUR
 ; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
 ; FILE REFERENCE: BET 03P0456
 ; CURRENT APPLICATION NUMBER: US/10/431,587
 ; CURRENT FILING DATE: 2003-05-08
 ; PRIOR APPLICATION NUMBER: FR 0205808
 ; PRIOR FILING DATE: 2002-05-10
 ; NUMBER OF SEQ ID NOS: 33
 ; SOFTWARE: PatentIn version 3.1
 ; SEQ ID NO 5
 ; LENGTH: 55
 ; TYPE: PRT
 ; ORGANISM: Hepatitis C virus
 ; US-10-431-587-5

Query Match 93.4%; Score 99; DB 12; Length 75;
 Best Local Similarity 95.0%; Pred. No. 1.e-06; 0; Gaps 0;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

RESULT 22
 US-10-431-587-16
 ; Sequence 16, Application US/10431587
 ; Publication No. US20040072267A1
 ; GENERAL INFORMATION:
 ; APPLICANT: BIORAD PASTEUR
 ; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
 ; FILE REFERENCE: BET 03P0456
 ; CURRENT APPLICATION NUMBER: US/10/431,587
 ; CURRENT FILING DATE: 2003-05-08
 ; PRIOR APPLICATION NUMBER: FR 0205808
 ; PRIOR FILING DATE: 2002-05-10
 ; NUMBER OF SEQ ID NOS: 33
 ; SOFTWARE: PatentIn version 3.1
 ; SEQ ID NO 16
 ; LENGTH: 75
 ; TYPE: PRT
 ; ORGANISM: Hepatitis C virus
 ; US-10-431-587-16

Query Match 93.4%; Score 99; DB 12; Length 75;
 Best Local Similarity 95.0%; Pred. No. 1.e-06; 0; Gaps 0;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

RESULT 23
 US-10-431-587-17
 ; Sequence 17, Application US/10431587
 ; Publication No. US20040072267A1
 ; GENERAL INFORMATION:
 ; APPLICANT: BIORAD PASTEUR
 ; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
 ; FILE REFERENCE: BET 03P0456
 ; CURRENT APPLICATION NUMBER: US/10/431,587
 ; CURRENT FILING DATE: 2003-05-08
 ; PRIOR APPLICATION NUMBER: FR 0205808
 ; PRIOR FILING DATE: 2002-05-10
 ; NUMBER OF SEQ ID NOS: 33
 ; SOFTWARE: PatentIn version 3.1
 ; SEQ ID NO 17
 ; LENGTH: 75
 ; TYPE: PRT
 ; ORGANISM: Hepatitis C virus

RESULT 21
 US-10-431-587-14
 ; Sequence 14, Application US/10431587
 ; Publication No. US20040072267A1
 ; GENERAL INFORMATION:
 ; APPLICANT: BIORAD PASTEUR
 ; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
 ; FILE REFERENCE: BET 03P0456
 ; CURRENT APPLICATION NUMBER: US/10/431,587
 ; CURRENT FILING DATE: 2003-05-08
 ; PRIOR APPLICATION NUMBER: FR 0205808
 ; PRIOR FILING DATE: 2002-05-10
 ; NUMBER OF SEQ ID NOS: 33

US-10-431-587-17

Query Match 93.4%; Score 99; DB 12; Length 75;
Best Local Similarity 95.0%; Pred. No. 1.4e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRTKTKNTNRRPQ 20
Db 1 MSTNPKPQRTKTKNTNRRPQ 20

RESULT 24
US-10-431-587-18
Sequence 18, Application US/10431587
Publication No. US20040072267A1

GENERAL INFORMATION:
APPLICANT: BIORAD PASTEUR
TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism

TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism

CURRENT APPLICATION NUMBER: US/10/431,587
CURRENT FILING DATE: 2003-05-08
PRIOR APPLICATION NUMBER: FR 0205808
PRIOR FILING DATE: 2002-05-10
NUMBER OF SEQ ID NOS: 33
SEQ ID NO: 18
LENGTH: 75
TYPE: PRT
ORGANISM: Hepatitis C virus
FEATURE:
NAME/KEY: MISC_FEATURE
LOCATION: (35) . (35)
OTHER INFORMATION: homo-serine

US-10-431-587-18

Query Match 93.4%; Score 99; DB 12; Length 75;
Best Local Similarity 95.0%; Pred. No. 1.4e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRTKTKNTNRRPQ 20
Db 1 MSTNPKPQRTKTKNTNRRPQ 20

RESULT 25
US-10-431-587-15
Sequence 15, Application US/10431587
Publication No. US20040072267A1

GENERAL INFORMATION:
APPLICANT: BIORAD PASTEUR
TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism

TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism

CURRENT APPLICATION NUMBER: US/10/431,587
CURRENT FILING DATE: 2003-05-08
PRIOR APPLICATION NUMBER: FR 0205808
PRIOR FILING DATE: 2002-05-10
NUMBER OF SEQ ID NOS: 33
SEQ ID NO: 15
LENGTH: 76
TYPE: PRT
ORGANISM: Hepatitis C virus

US-10-431-587-15

Query Match 93.4%; Score 99; DB 12; Length 76;
Best Local Similarity 95.0%; Pred. No. 1.5e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRTKTKNTNRRPQ 20
Db 1 MSTNPKPQRTKTKNTNRRPQ 20

RESULT 26
US-09-758-308-1
Sequence 1, Application US/09758308
Patent No. US20020090607A1

GENERAL INFORMATION:
APPLICANT: HOWARD A. FIELDS AND YURY B. KHUDYAKOV
TITLE OF INVENTION: ANTIGENIC EPITOPEs AND MOSAIC POLYPEPTIDES OF HEPATITIS C VIRUS
TITLE OF INVENTION: PROTEINS
FILE REFERENCE: 14114_0349U2
CURRENT APPLICATION NUMBER: US/09/758,308
CURRENT FILING DATE: 2001-01-10
PRIOR APPLICATION NUMBER: 60/092,339
PRIOR FILING DATE: 1999-07-10
NUMBER OF SEQ ID NOS: 5
SOFTWARE: PatentIn version 3.0
SEQ ID NO: 1
TYPE: PRT
ORGANISM: Hepatitis C Virus

US-09-758-308-1

Query Match 93.4%; Score 99; DB 9; Length 91;
Best Local Similarity 95.0%; Pred. No. 1.7e-06;
Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRTKTKNTNRRPQ 20
Db 1 MSTNPKPQRTKTKNTNRRPQ 20

RESULT 27
US-09-756-875-8
Sequence 8, Application US/09756875
Patent No. US20020150990A1

GENERAL INFORMATION:
APPLICANT: PIKE, IAN
TITLE OF INVENTION: HEPATITIS C VIRUS PEPTIDES
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Suite 701-E Columbia Square
STREET: 555 13th Street, N. W.
CITY: Washington
STATE: D. C.
COUNTRY: U. S.
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/756,875
FILING DATE:
CLASSIFICATION:
PRIORITY APPLICATION DATA:
APPLICATION NUMBER: US 08/259,721
FILING DATE: 29-AUG-1994
APPLICATION NUMBER: PCT/GB93/00410
FILING DATE: 26-FEB-1993
ATTORNEY/AGENT INFORMATION:
NAME: ERNST, BARBARA G.
REGISTRATION NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 1808-157A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 781-6040
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 97 amino acids
TYPE: amino acid
TOPOLOGY: unknown
MOLECULE TYPE: peptide

US-09-756-875-8
 Query Match 93.4%; Score 99; DB 9; Length 97;
 Best Local Similarity 95.0%; Pred. No. 1.e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKQTKTRNTNRRQ 20
 Db 1 MSTNPKPQTKTRNTNRRQ 20

RESULT 28

; Sequence 77, Application US/09921397
 ; Patent No. US20020151484A1

GENERAL INFORMATION:

; APPLICANT: HYBRIDGENICS

; TITLE OF INVENTION: S1D nucleic acids and polypeptides selected from a pathogenic strain of the hepatitis C virus and applications thereof

; FILE REFERENCE: B4809A - JAZ

CURRENT APPLICATION NUMBER: US/09/921,397

CURRENT FILING DATE: 2001-08-02

PRIOR APPLICATION NUMBER: EP 00402225.7

PRIOR FILING DATE: 2000-08-03

NUMBER OF SEQ ID NOs: 156

SEQ ID NO 77

LENGTH: 103

TYPE: PCT

ORGANISM: Hepatitis C virus

US 09-221-397-77

Query Match 93.4%; Score 99; DB 9; Length 103;
 Best Local Similarity 95.0%; Pred. No. 2.e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKQTKTRNTNRRQ 20
 Db 1 MSTNPKPQTKTRNTNRRQ 33

RESULT 29

; Sequence 14, Application US/09851138
 ; PUBLICACION NO. US20020183508A1

GENERAL INFORMATION:

; APPLICANT: MAERTENS, GERT

; STUYVER, LIEVEN
 TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS

NUMBER OF SEQUENCES: 207

CORRESPONDENCE ADDRESS:

ADDRESSEE: ARNOLD, WHITE & DURKEE
 STREET: P.O. BOX 4433
 CITY: HOUSTON
 STATE: TEXAS
 COUNTRY: USA

ZIP: 77210-4433

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/851,138

FILING DATE: 09-May-2001

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/836,075

FILING DATE: <Unknown>

APPLICATION NUMBER: EP 94870166.9

FILING DATE: 21 Oct 1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/836,075

FILING DATE: <Unknown>

APPLICATION NUMBER: EP 94870166.9

FILING DATE: 21 Oct 1994

APPLICATION NUMBER: EP 95870076.7

FILING DATE: 28 Jun 1995

ATTORNEY/AGENT INFORMATION:

; NAME: KAMMERER, PATRICIA A.

; REGISTRATION NUMBER: 29,775

; REFERENCE/DOCKET NUMBER: INNS:004

INFORMATION FOR SEQ ID NO: 14:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 106 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: peptide

; SEQUENCE DESCRIPTION: SEQ ID NO: 14:

US-09-851-138-14

Query Match 93.4%; Score 99; DB 9; Length 108;

; Best Local Similarity 95.0%; Pred. No. 2.1e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPKQTKTRNTNRRQ 20

Db 1 MSTNPKPQTKTRNTNRRQ 20

RESULT 30

US-09-851-138-6

; Sequence 6, Application US/09851138

; Publication No. US20020183508A1

GENERAL INFORMATION:

; APPLICANT: MAERTENS, GERT

; STUYVER, LIEVEN

TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS

NUMBER OF SEQUENCES: 207

CORRESPONDENCE ADDRESS:

ADDRESSEE: ARNOLD, WHITE & DURKEE
 STREET: P.O. BOX 4433
 CITY: HOUSTON
 STATE: TEXAS
 COUNTRY: USA

ZIP: 77210-4433

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/851,138

FILING DATE: 09-May-2001

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/836,075

FILING DATE: <Unknown>

APPLICATION NUMBER: EP 94870166.9

FILING DATE: 21 Oct 1994

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/836,075

FILING DATE: <Unknown>

APPLICATION NUMBER: EP 94870166.9

FILING DATE: 21 Oct 1994

ATTORNEY/AGENT INFORMATION:

; NAME: KAMMERER, PATRICIA A.

; REGISTRATION NUMBER: 29,775

; REFERENCE/DOCKET NUMBER: INNS:004

INFORMATION FOR SEQ ID NO: 1:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 109 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: peptide

; SEQUENCE DESCRIPTION: SEQ ID NO: 1:

US-09-851-138-6

Query Match 93.4%; Score 99; DB 9; Length 109;

; Best Local Similarity 95.0%; Pred. No. 2.1e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

REGISTRATION NUMBER: 24,618
 REFERENCE DOCKET NUMBER: 2084-033-0
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (703) 413-3000
 TELEFAX: (703) 413-2220
 INFORMATION FOR SEQ ID NO: 4;
 SEQUENCE CHARACTERISTICS:
 LENGTH: 120 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 SEQUENCE DESCRIPTION: SEQ ID NO: 4:
 US-09-306-780-4

Query Match 93.4%; Score 99; DB 12; Length 120;
 Best Local Similarity 95.0%; Pred. No. 2.5e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPORKTKNTNRRPQ 20
 Db 1 MSTNPKPQPKTKNTNRRPQ 20

RESULT 33
 US-10-268-569-19
 Sequence 19, Application US/10268569
 Publication No US20030152965A1
 GENERAL INFORMATION:
 APPLICANT: Ortho-Clinical Diagnostics, Inc.
 TITLE OF INVENTION: HCV Core Protein Sequences
 FILE REFERENCE: CDS-088
 CURRENT APPLICATION NUMBER: US20030152965A1
 PRIORITY DATE: 2002-10-10
 PRIOR APPLICATION NUMBER: 60/347,303
 NUMBER OF SEQ ID NOS: 19
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO 19
 LENGTH: 130
 TYPE: PRT
 ORGANISM: Hepatitis C virus

US-10-268-569-19

Query Match 93.4%; Score 99; DB 14; Length 130;
 Best Local Similarity 95.0%; Pred. No. 2.5e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPORKTKNTNRRPQ 20
 Db 1 MSTNPKPQPKTKNTNRRPQ 20

RESULT 32
 US-09-306-780-4
 Sequence 4, Application US/09306780
 PUBLIC "-//IETF//DTD HTML 2.0//EN"
 GENERAL INFORMATION:
 APPLICANT: TAKEMURA, FUMINORI
 ITOH, SATOHI
 ITOH, EIICHI
 TITLE OF INVENTION: NUCLEIC ACID-BOUND POLYPEPTIDE, METHOD
 OF PRODUCING NUCLEIC ACID-BOUND POLYPEPTIDE AND
 NUMBER OF SEQUENCES: 20
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: OBLON, SPIVAK, McCLELLAND, MAIER & NEUSTADT,
 P.C.
 STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: U.S.A.
 ZIP: 22202
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/306,780
 FILING DATE: 07-May-1999
 CLASSIFICATION: <Unknown>
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US/08/841,657A
 FILING DATE: 30-Apr-1997
 APPLICATION NUMBER: JP 8-134444
 FILING DATE: 01-MAY-1997
 ATTORNEY/AGENT INFORMATION:
 NAME: OBLON, NORMAN F.

RESULT 34
 US-09-051-138-60
 Sequence 60, Application US/09851138
 Publication No. US20020163508A1
 GENERAL INFORMATION:
 APPLICANT: MAERTENS, GERT
 STOYVER, LIEVEN
 TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES
 AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC
 AGENTS
 NUMBER OF SEQUENCES: 207
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: ARNOLD, WHITE & DURKEE
 STREET: P.O. BOX 4433
 CITY: HOUSTON
 STATE: TEXAS
 COUNTRY: USA
 ZIP: 77210-4433
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/306,780
 FILING DATE: 07-May-1999
 CLASSIFICATION: <Unknown>
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US/08/841,657A
 FILING DATE: 30-Apr-1997
 APPLICATION NUMBER: JP 8-134444
 FILING DATE: 01-MAY-1997
 ATTORNEY/AGENT INFORMATION:
 NAME: OBLON, NORMAN F.

OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Microsoft Word 6.0 / ASCII text output
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/851,138
 FILING DATE: 09-MAY-2001
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: PCT/US01/051336A1
 FILING DATE: <Unknown>
 ATTORNEY/AGENT INFORMATION:
 NAME: KAMMERER, PATRICIA A.
 REGISTRATION NUMBER: 29,775
 INFORMATION FOR SEQ ID NO: 60:
 REFERENCE CHARACTERISTICS:
 SEQUENCE DESCRIPTION:
 LENGTH: 138 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 SEQUENCE DESCRIPTION: SEQ ID NO: 60:
 US-09-851-138-60

Query Match 93.4%; Score 99; DB 9; Length 138;
 Best Local Similarity 95.0%; Pred. No. 2.6e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPORTKTKRNTNRRPQ 20
 Db 1 MSTNPKPQRTKTKRNTNRRPQ 20

RESULT 35
 US-10-292-129-14
 Sequence 14, Application US/10292129
 Publication No. US20030148267A1
 GENERAL INFORMATION:
 APPLICANT: Schmidt, Emmett Vance
 TITLE OF INVENTION: SCREENING ASSAY FOR HEPATITIS C VIRUS
 FILE REFERENCE: 00786-539001
 CURRENT APPLICATION NUMBER: US/10/292,129
 CURRENT FILING DATE: 2002-11-08
 PRIOR APPLICATION NUMBER: US 60/345,405
 PRIOR FILING DATE: 2001-11-09
 SOFTWARE: FastSEQ for Windows Version 4.0
 SEQ ID NO 14
 LENGTH: 151
 TYPE: PPT
 ORGANISM: Hepatitis C virus
 US-10-292-129-14

Query Match 93.4%; Score 99; DB 14; Length 151;
 Best Local Similarity 95.0%; Pred. No. 2.6e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPORTKTKRNTNRRPQ 20
 Db 1 MSTNPKPQRTKTKRNTNRRPQ 20

RESULT 36
 US-09-306-780-8
 Sequence 8, Application US/09306780
 Publication No. US2010051336A1
 GENERAL INFORMATION:
 APPLICANT: TAKEMURA, FUMINORI
 UENO, EIICHI
 ITOH, SATORU

TITLE OF INVENTION: NUCLEIC ACID-BOUND POLYPEPTIDE, METHOD OF PRODUCING NUCLEIC ACID-BOUND POLYPEPTIDE AND IMMUNOASSAY USING THE POLYPEPTIDE.
 NUMBER OF SEQUENCES: 20
 CORRESPONDENCE ADDRESS: P.O. BOX 22202, OBLON, SPIVAK, McCLELLAND, MAIER & NEUSTADT, P.C.
 STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: U.S.A.
 ZIP: 22202
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0., Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/306,780
 FILING DATE: 07-May-1999
 CLASSIFICATION: <Unknown>
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US/08/841,657A
 FILING DATE: 30-APR-1997
 APPLICATION NUMBER: JP 8-134444
 FILING DATE: 01-MAY-1997
 ATTORNEY/AGENT INFORMATION:
 NAME: OBLON, NORMAN F.
 REGISTRATION NUMBER: 24,618
 REFERENCE/DOCKET NUMBER: 2084-033-0
 TELEPHONE: (703) 413-3000
 TELEFAX: (703) 413-2220
 INFORMATION FOR SEQ ID NO: 8:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 161 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 SEQUENCE DESCRIPTION: SEQ ID NO: 8:
 US-09-306-780-8

Query Match 93.4%; Score 99; DB 12; Length 161;
 Best Local Similarity 95.0%; Pred. No. 3e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MSTIPKPORTKTKRNTNRRPQ 20
 Db 6 MSTNPKPQRTKTKRNTNRRPQ 25

RESULT 37
 US-09-899-046-152
 Sequence 152, Application US/09899046
 Publication No. US20030008274A1
 GENERAL INFORMATION:
 APPLICANT:
 TITLE OF INVENTION: New sequences of hepatitis C virus
 NUMBER OF SEQUENCES: 270
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0., Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/899,046
 FILING DATE:
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: 08/362,455
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 152:
 SEQUENCE CHARACTERISTICS:

LENGTH: 166 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-09-859-046-152

Query Match 93.4%; Score 99; DB 10; Length 166;
 Best Local Similarity 95.0%; Pred. No. 3.1e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTNRRPQ 20
 Db 1 MSTNPKPQRTKRTNRRPQ 20

RESULT 38
 US-09-878-281-152
 Sequence 152, Application US/09878281
 Publication No. US20030032005A1

GENERAL INFORMATION:
 APPLICANT:
 TITLE OF INVENTION: New sequences of hepatitis C virus
 NUMBER OF SEQUENCES: 270
 COMPUTER READABLE FORM:
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/878,281
 FILING DATE:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/362,455
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 152:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 166 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-09-878-281-152

Query Match 93.4%; Score 99; DB 10; Length 166;
 Best Local Similarity 95.0%; Pred. No. 3.1e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTNRRPQ 20
 Db 1 MSTNPKPQRTKRTNRRPQ 20

RESULT 39
 US-09-873-224-152
 Sequence 152, Application US/09873224
 Publication No. US20030064360A1

GENERAL INFORMATION:
 APPLICANT: <Unknown>
 TITLE OF INVENTION: New sequences of hepatitis C virus
 NUMBER OF SEQUENCES: 270
 COMPUTER READABLE FORM:
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)

SEARCH RESULTS:
 Search completed: July 20, 2004, 09:45:29
 Job time : 41 secs

FILING DATE: 05-Jun-2001
 CLASSIFICATION: <Unknown>
 PRIORITY APPLICATION NUMBER: 08/362,455
 ATTORNEY/AGENT INFORMATION:
 NAME: Immunetics S.
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 00 32 9 241 07 11
 TELEFAX: 00 32 9 241 07 99
 INFORMATION FOR SEQ ID NO: 152:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 169 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 SEQUENCE DESCRIPTION: SEQ ID NO: 152:

US-09-873-224-152

Query Match 93.4%; Score 99; DB 12; Length 166;
 Best Local Similarity 95.0%; Pred. No. 3.1e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTNRRPQ 20
 Db 1 MSTNPKPQRTKRTNRRPQ 20

RESULT 40
 US-09-899-046-42
 Sequence 42, Application US/09899046
 Publication No. US2003008274A1

GENERAL INFORMATION:
 APPLICANT:
 TITLE OF INVENTION: New sequences of hepatitis C virus
 NUMBER OF SEQUENCES: 270
 COMPUTER READABLE FORM:
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/899,046
 FILING DATE:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/362,455
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 42:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 169 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-09-899-046-42

Query Match 93.4%; Score 99; DB 10; Length 169;
 Best Local Similarity 95.0%; Pred. No. 3.2e-06;
 Matches 19; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MST1PKPQRTKRTNRRPQ 20
 Db 1 MSTNPKPQRTKRTNRRPQ 20

Search completed: July 20, 2004, 09:45:29
 Job time : 41 secs

APPLICATION NUMBER: US/09/873,224

ALIGNMENTS

TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
ANTIBODIES TO HEPATITIS C VIRUS

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:
ADDRESSEES: NIXON & VANDEHYDE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
STATE: VA
COUNTRY: USA
ZIP: 22201

COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/941,611
FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/391,671
FILING DATE: 1-95-02-21
APPLICATION NUMBER: WO PCT/EP91/02409
FILING DATE: 13-DEC-1991
APPLICATION NUMBER: EP 90124241.2
FILING DATE: 14-DEC-1990

ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
DEPARTMENT: NY/DO

SUMMARIES

Result No.	Score	Query	Length	DB	ID	Description	
						\$	%
1	110	100.0	20	9	US-09-941-611-2	Sequence 2,	Appli
2	110	100.0	20	14	US-10-044-955-2	Sequence 2,	Appli
3	110	100.0	30	12	US-10-296-734-488	Sequence 8,	Appli
4	110	100.0	44	14	US-10-367-677-1	Sequence 1,	Appli
5	110	100.0	48	9	US-09-851-138-22	Sequence 22,	Appli
6	110	100.0	53	12	US-10-431-587-3	Sequence 3,	Appli
7	110	100.0	53	12	US-10-431-587-10	Sequence 10,	Appli
8	110	100.0	53	12	US-10-431-587-11	Sequence 11,	Appli
9	110	100.0	53	12	US-10-431-587-12	Sequence 12,	Appli
10	110	100.0	63	12	US-10-431-587-2	Sequence 2,	Appli
11	110	100.0	63	12	US-10-431-587-7	Sequence 7,	Appli
12	110	100.0	63	12	US-10-431-587-8	Sequence 8,	Appli
13	110	100.0	70	12	US-10-431-587-9	Sequence 9,	Appli
14	110	100.0	73	12	US-10-431-587-13	Sequence 13,	Appli
15	110	100.0	74	12	US-10-431-587-14	Sequence 14,	Appli
16	110	100.0	74	12	US-10-431-587-15	Sequence 15,	Appli
17	110	100.0	74	12	US-10-431-587-16	Sequence 16,	Appli
18	110	100.0	74	12	US-10-431-587-17	Sequence 17,	Appli
19	110	100.0	74	12	US-10-431-587-18	Sequence 18,	Appli
20	110	100.0	74	12	US-10-431-587-19	Sequence 19,	Appli
21	110	100.0	74	12	US-10-431-587-20	Sequence 20,	Appli
22	110	100.0	74	12	US-10-431-587-21	Sequence 21,	Appli
23	110	100.0	74	12	US-10-431-587-22	Sequence 22,	Appli
24	110	100.0	74	12	US-10-431-587-23	Sequence 23,	Appli
25	110	100.0	74	12	US-10-431-587-24	Sequence 24,	Appli
26	110	100.0	74	12	US-10-431-587-25	Sequence 25,	Appli
27	110	100.0	74	12	US-10-431-587-26	Sequence 26,	Appli
28	110	100.0	74	12	US-10-431-587-27	Sequence 27,	Appli
29	110	100.0	74	12	US-10-431-587-28	Sequence 28,	Appli
30	110	100.0	74	12	US-10-431-587-29	Sequence 29,	Appli
31	110	100.0	74	12	US-10-431-587-30	Sequence 30,	Appli
32	110	100.0	74	12	US-10-431-587-31	Sequence 31,	Appli
33	110	100.0	74	12	US-10-431-587-32	Sequence 32,	Appli
34	110	100.0	74	12	US-10-431-587-33	Sequence 33,	Appli
35	110	100.0	74	12	US-10-431-587-34	Sequence 34,	Appli
36	110	100.0	74	12	US-10-431-587-35	Sequence 35,	Appli
37	110	100.0	74	12	US-10-431-587-36	Sequence 36,	Appli
38	110	100.0	74	12	US-10-431-587-37	Sequence 37,	Appli
39	110	100.0	74	12	US-10-431-587-38	Sequence 38,	Appli
40	110	100.0	74	12	US-10-431-587-39	Sequence 39,	Appli
41	110	100.0	74	12	US-10-431-587-40	Sequence 40,	Appli
42	110	100.0	74	12	US-10-431-587-41	Sequence 41,	Appli
43	110	100.0	74	12	US-10-431-587-42	Sequence 42,	Appli
44	110	100.0	74	12	US-10-431-587-43	Sequence 43,	Appli
45	110	100.0	74	12	US-10-431-587-44	Sequence 44,	Appli
46	110	100.0	74	12	US-10-431-587-45	Sequence 45,	Appli
47	110	100.0	74	12	US-10-431-587-46	Sequence 46,	Appli
48	110	100.0	74	12	US-10-431-587-47	Sequence 47,	Appli
49	110	100.0	74	12	US-10-431-587-48	Sequence 48,	Appli
50	110	100.0	74	12	US-10-431-587-49	Sequence 49,	Appli
51	110	100.0	74	12	US-10-431-587-50	Sequence 50,	Appli
52	110	100.0	74	12	US-10-431-587-51	Sequence 51,	Appli
53	110	100.0	74	12	US-10-431-587-52	Sequence 52,	Appli
54	110	100.0	74	12	US-10-431-587-53	Sequence 53,	Appli
55	110	100.0	74	12	US-10-431-587-54	Sequence 54,	Appli
56	110	100.0	74	12	US-10-431-587-55	Sequence 55,	Appli
57	110	100.0	74	12	US-10-431-587-56	Sequence 56,	Appli
58	110	100.0	74	12	US-10-431-587-57	Sequence 57,	Appli
59	110	100.0	74	12	US-10-431-587-58	Sequence 58,	Appli
60	110	100.0	74	12	US-10-431-587-59	Sequence 59,	Appli
61	110	100.0	74	12	US-10-431-587-60	Sequence 60,	Appli
62	110	100.0	74	12	US-10-431-587-61	Sequence 61,	Appli
63	110	100.0	74	12	US-10-431-587-62	Sequence 62,	Appli
64	110	100.0	74	12	US-10-431-587-63	Sequence 63,	Appli
65	110	100.0	74	12	US-10-431-587-64	Sequence 64,	Appli
66	110	100.0	74	12	US-10-431-587-65	Sequence 65,	Appli
67	110	100.0	74	12	US-10-431-587-66	Sequence 66,	Appli
68	110	100.0	74	12	US-10-431-587-67	Sequence 67,	Appli
69	110	100.0	74	12	US-10-431-587-68	Sequence 68,	Appli
70	110	100.0	74	12	US-10-431-587-69	Sequence 69,	Appli
71	110	100.0	74	12	US-10-431-587-70	Sequence 70,	Appli
72	110	100.0	74	12	US-10-431-587-71	Sequence 71,	Appli
73	110	100.0	74	12	US-10-431-587-72	Sequence 72,	Appli
74	110	100.0	74	12	US-10-431-587-73	Sequence 73,	Appli
75	110	100.0	74	12	US-10-431-587-74	Sequence 74,	Appli
76	110	100.0	74	12	US-10-431-587-75	Sequence 75,	Appli
77	110	100.0	74	12	US-10-431-587-76	Sequence 76,	Appli
78	110	100.0	74	12	US-10-431-587-77	Sequence 77,	Appli
79	110	100.0	74	12	US-10-431-587-78	Sequence 78,	Appli
80	110	100.0	74	12	US-10-431-587-79	Sequence 79,	Appli
81	110	100.0	74	12	US-10-431-587-80	Sequence 80,	Appli
82	110	100.0	74	12	US-10-431-587-81	Sequence 81,	Appli
83	110	100.0	74	12	US-10-431-587-82	Sequence 82,	Appli
84	110	100.0	74	12	US-10-431-587-83	Sequence 83,	Appli
85	110	100.0	74	12	US-10-431-587-84	Sequence 84,	Appli
86	110	100.0	74	12	US-10-431-587-85	Sequence 85,	Appli
87	110	100.0	74	12	US-10-431-587-86	Sequence 86,	Appli
88	110	100.0	74	12	US-10-431-587-87	Sequence 87,	Appli
89	110	100.0	74	12	US-10-431-587-88	Sequence 88,	Appli
90	110	100.0	74	12	US-10-431-587-89	Sequence 89,	Appli
91	110	100.0	74	12	US-10-431-587-90	Sequence 90,	Appli
92	110	100.0	74	12	US-10-431-587-91	Sequence 91,	Appli
93	110	100.0	74	12	US-10-431-587-92	Sequence 92,	Appli
94	110	100.0	74	12	US-10-431-587-93	Sequence 93,	Appli
95	110	100.0	74	12	US-10-431-587-94	Sequence 94,	Appli
96	110	100.0	74	12	US-10-431-587-95	Sequence 95,	Appli
97	110	100.0	74	12	US-10-431-587-96	Sequence 96,	Appli
98	110	100.0	74	12	US-10-431-587-97	Sequence 97,	Appli
99	110	100.0	74	12	US-10-431-587-98	Sequence 98,	Appli
100	110	100.0	74	12	US-10-431-587-99	Sequence 99,	Appli
101	110	100.0	74	12	US-10-431-587-100	Sequence 100,	Appli

REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 SEQUENCE DESCRIPTION: SEQ ID NO: 2:
 US-09-941 611-2

Query Match 100.0%; Score 110; DB 9; Length 20;
 Best Local Similarity 100.0%; Pred. No. 9e-10; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 1 PQRKTKRNTNRRPQDVKFPG 20

RESULT 2
 US-10-044-995-2
 ; Sequence 2, Application US/10044995
 ; Publication No. US20010049285A1
 GENERAL INFORMATION:
 APPLICANT: DELLEYS, ROBERT J
 POLELET, DIRK
 MAETTEENS, GERT
 VAN HEUVERSWYN, HUGO

TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 ZIP: 22201

COMPUTER READABLE FORM:
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/10/044,995
 FILING DATE: 15-Jan-2002
 CLASSIFICATION: <Unknown>
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/391,671
 FILING DATE: <Unknown>
 APPLICATION NUMBER: US 07/920,286
 FILING DATE: 14-OCT-1992
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 13-DEC-1991
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SAOOFF, B.J.
 REGISTRATION NUMBER: 36,663
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 TELEX: 70381641000

INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear

; MOLECULE TYPE: peptide
 ; SEQUENCE DESCRIPTION: SEQ ID NO: 2:
 ; US-10-044-995-2
 Query Match 100.0%; Score 110; DB 14; Length 20;
 Best Local Similarity 100.0%; Pred. No. 9e-10; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 1 PQRKTKRNTNRRPQDVKFPG 20

RESULT 3
 US-10-296-734-408
 ; Sequence 408, Application US/10296734
 ; Publication No. US20040054137A1
 GENERAL INFORMATION:
 APPLICANT: Thompson, Scott A
 ; TITLE OF INVENTION: Synthetic molecules and uses therefor
 ; FILE REFERENCE: Savine
 CURRENT APPLICATION NUMBER: US/10/296,734
 ; CURRENT FILING DATE: 2003-08-04
 ; PRIOR APPLICATION NUMBER: AU PQ7761/00
 ; PRIOR FILING DATE: 2000-05-26
 ; NUMBER OF SEQ ID NOS: 1507
 ; SOFTWARE: PatentIn version 3.2
 ; SEQ ID NO: 408
 ; LENGTH: 30
 ; TYPE: PRT
 ; ORGANISM: Artificial
 ; FEATURE:
 ; OTHER INFORMATION: HepC 1a segment 1

US-10-296-734-408

Query Match 100.0%; Score 110; DB 12; Length 30;
 Best Local Similarity 100.0%; Pred. No. 1.4e-09; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
 Db 9 PQRKTKRNTNRRPQDVKFPG 28

RESULT 4
 US-10-367-677-1
 ; Sequence 1, Application US/10367677
 ; Publication No. US20030118604A1
 GENERAL INFORMATION:
 ; APPLICANT: JOLIVET, MICHEL
 ; FILE REFERENCE: 103959
 ; APPENDANT: PENIN, FRANCOIS
 ; APPENDANT: DALBON, PASCAL
 ; APPENDANT: LADAVIERE, LAURENT
 ; APPENDANT: LACOUX, XAVIER
 ; TITLE OF INVENTION: TREATING AN HCV INFECTION
 ; CURRENT APPLICATION NUMBER: US/10/367,677
 ; CURRENT FILING DATE: 2003-02-19
 ; PRIOR APPLICATION NUMBER: US/09/389,756
 ; PRIOR FILING DATE: 1999-09-07.
 ; PRIOR APPLICATION NUMBER: earlier application number: PCT/ER98/00442
 ; PRIOR FILING DATE: earlier filing date: 1998-03-05
 ; NUMBER OF SEQ ID NOS: 11
 ; SOFTWARE: PatentIn Ver. 2.1
 ; SEQ ID NO: 1
 ; LENGTH: 44
 ; TYPE: PRT
 ; ORGANISM: Hepatitis C virus
 ; PUBLICATION INFORMATION:
 ; AUTHORS: Ogata, N. et al.

RESULT 4
US-10-431-587-3
; Sequence 3, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; FILE REFERENCE: BET 03P456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 02056008
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-3

Query Match Score 110; DB 14; Length 4;
Best Local Similarity 100.0%; Pred. No. 2.1e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 6 PQRKTKRNTNRRPQDVKFPG 25

RESULT 5
US-09-851-138-22
; Sequence 22, Application US/098511138
; Publication No. US20020183508A1
; GENERAL INFORMATION:
; APPLICANT: MARTEENS, GERT
; STOYVER, LIEVEN
; TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS
; NUMBER OF SEQUENCES: 207
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ARNOLD, WHITE & DURKEE
; STREET: P.O. BOX 4433
; CITY: HOUSTON
; STATE: TEXAS
; ZIP: 77210-4433
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Microsoft Word 6.0 / ASCII text output
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/851,138
; FILING DATE: 09-May-2001
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/836,075
; FILING DATE: <Unknown>
; APPLICATION NUMBER: EP 94870166.9
; FILING DATE: 21 Oct 1994
; APPLICATION NUMBER: EP 95870076.7
; FILING DATE: 28 Jun 1995
; ATTORNEY/AGENT INFORMATION:
; NAME: KAMMERER, PATRICIA A.
; REGISTRATION NUMBER: 29,775
; REFERENCE/DOCKET NUMBER: INNS: 004
; INFORMATION FOR SEQ ID NO: 22;
; SEQUENCE CHARACTERISTICS:
; LENGTH: 48 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: Peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 22;
; US-09-851-138-22

Query Match Score 110; DB 9; Length 48;
Best Local Similarity 100.0%; Pred. No. 2.3e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 6
US-10-431-587-3
; Sequence 10, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; FILE REFERENCE: BET 03P456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 02056008
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-3

Query Match Score 110; DB 12; Length 53;
Best Local Similarity 100.0%; Pred. No. 2.5e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 7
US-10-431-587-10
; Sequence 10, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; FILE REFERENCE: BET 03P456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 02056008
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 10
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-10

Query Match Score 110; DB 12; Length 53;
Best Local Similarity 100.0%; Pred. No. 2.5e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 8
US-10-431-587-11
; Sequence 11, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; FILE REFERENCE: BET 03P456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 02056008
; PRIOR FILING DATE: 2002-05-10

NUMBER OF SEQ ID NOS: 33
; SEQ ID NO: 11
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-11

Query Match 100.0%; Score 110; DB 12; Length 53;
Best Local Similarity 100.0%; Pred. No. 2.5e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRTKRNTRRPPDVKPG 20
Db 7 PQRTKRNTRRPPDVKPG 26

RESULT 9
US-10-431-587-12
; Sequence 12, Application US/10431587
; Publication No. US20040072267A1
GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; CURRENT APPLICATION NUMBER: US/10/431.587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 12
; LENGTH: 53
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-12

Query Match 100.0%; Score 110; DB 12; Length 53;
Best Local Similarity 100.0%; Pred. No. 2.5e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRTKRNTRRPPDVKPG 20
Db 7 PQRTKRNTRRPPDVKPG 26

RESULT 10
US-10-431-587-2
; Sequence 2, Application US/10431587
; Publication No. US20040072267A1
GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; CURRENT APPLICATION NUMBER: US/10/431.587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 2
; LENGTH: 63
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-2

Query Match 100.0%; Score 110; DB 12; Length 63;
Best Local Similarity 100.0%; Pred. No. 3e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRTKRNTRRPPDVKPG 20

Db 2 PQRTKRNTRRPPDVKPG 21

RESULT 11
US-10-431-587-7
; Sequence 7, Application US/10431587
; Publication No. US20040072267A1
GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431.587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: PR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 7
; LENGTH: 63
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-7

Query Match 100.0%; Score 110; DB 12; Length 63;
Best Local Similarity 100.0%; Pred. No. 3e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRTKRNTRRPPDVKPG 20
Db 2 PQRTKRNTRRPPDVKPG 21

RESULT 12
US-10-431-587-8
; Sequence 8, Application US/10431587
; Publication No. US20040072267A1
GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431.587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: PR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 8
; LENGTH: 63
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-8

Query Match 100.0%; Score 110; DB 12; Length 63;
Best Local Similarity 100.0%; Pred. No. 3e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRTKRNTRRPPDVKPG 20
Db 2 PQRTKRNTRRPPDVKPG 21

RESULT 13
US-10-431-587-9
; Sequence 9, Application US/10431587
; Publication No. US20040072267A1
GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
; FILE REFERENCE: BET 03P0456

Qy 1 PQRTKRNTRRPPDVKPG 20

CURRENT APPLICATION NUMBER: US/10/431,587
 CURRENT FILING DATE: 2003-05-08
 PRIOR APPLICATION NUMBER: FR 0205808
 NUMBER OF SEQ ID NOS: 33
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 9
 LENGTH: 70
 TYPE: PRT
 ORGANISM: Hepatitis C virus
 US-10-431-587-9

Query Match 100.0%; Score 110; DB 12; Length 70;
 Best Local Similarity 100.0%; Pred. No. 3. 4e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 2 PQRKTKRNTNRRPQDVKEPG 21

RESULT 14
 US-10-431-587-13
 Sequence 13, Application US/10431587
 Publication No. US20040072267A1
 GENERAL INFORMATION:
 APPLICANT: BIORAD PASTEUR
 TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody against, an infectious microorganism
 FILE REFERENCE: BET 03P0156
 CURRENT APPLICATION NUMBER: US/10/431,587
 CURRENT FILING DATE: 2003-05-08
 PRIOR APPLICATION NUMBER: FR 0205808
 NUMBER OF SEQ ID NOS: 33
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 13
 LENGTH: 73
 TYPE: PRT
 ORGANISM: Hepatitis C virus
 US-10-431-587-13

Query Match 100.0%; Score 110; DB 12; Length 73;
 Best Local Similarity 100.0%; Pred. No. 3. 5e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 15
 US-09-851-138-10
 Sequence 10, Application US/098511138
 Publication No. US20020183508A1
 GENERAL INFORMATION:
 APPLICANT: STOEVER, LIBETEN, GERT
 TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS
 NUMBER OF SEQUENCES: 207
 CORRESPONDENCE ADDRESS:
 ADDRESS: ARNOLD, WHITE & DURKEE
 STREET: P.O. BOX 4433
 CITY: HOUSTON
 STATE: TEXAS
 COUNTRY: USA
 ZIP: 77200-4433
 COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Microsoft Word 6.0 / ASCII text output
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/851,138
 FILING DATE: 09-MAY-2001
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/836,075
 FILING DATE: <Unknown>
 APPLICATION NUMBER: EP 94 870166.9
 FILING DATE: 21 Oct 1994
 APPLICATION NUMBER: EP 95 870076.7
 FILING DATE: 28 Jun 1995
 ATTORNEY/AGENT INFORMATION:
 NAME: KAMMERER, PATRICIA A.
 REGISTRATION NUMBER: 29,775
 REFERENCE/DOCKET NUMBER: INNS: 004
 INFORMATION FOR SEQ ID NO: 10:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 74 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 SEQUENCE DESCRIPTION: SEQ ID NO: 10:
 US-09-851-138-10

Query Match 100.0%; Score 110; DB 9; Length 74;
 Best Local Similarity 100.0%; Pred. No. 3. 6e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 16
 US-10-651-165-198
 Sequence 198, Application US/10651165
 Publication No. US20040047877A1
 GENERAL INFORMATION:
 APPLICANT: LEBOUX-ROELS, Geert
 APPLICANT: DELEYS, Robert
 APPLICANT: MAERTENS, Geert
 TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C
 FILE REFERENCE: 2551-94
 CURRENT APPLICATION NUMBER: US/10/651,165
 CURRENT FILING DATE: 2003-03-02
 PRIOR APPLICATION NUMBER: US/08/974,690C
 PRIOR FILING DATE: 1997-11-19
 PRIOR APPLICATION NUMBER: PCT/EP94/03555
 PRIOR FILING DATE: 1994-10-28
 PRIOR APPLICATION NUMBER: EP 93402718.6
 PRIOR FILING DATE: 1993-11-04
 NUMBER OF SEQ ID NOS: 266
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 198
 LENGTH: 74
 TYPE: PRT
 ORGANISM: hepatitis C virus
 US-10-651-165-198

Query Match 100.0%; Score 110; DB 12; Length 74;
 Best Local Similarity 100.0%; Pred. No. 3. 6e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
 Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 17
 US-10-131-587-1
 Sequence 1, Application US/10431587
 Publication No. US2004072267A1

```

; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 75
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-1
Query Match 100.0%; Score 110; DB 12; Length 75;
Best Local Similarity 100.0%; Pred. No. 3.6e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 PQRTKRNTRRQDVKFPG 20
Db 7 PQRTKRNTRRQDVKFPG 26

RESULT 20
US-10-431-587-6
; Sequence 6, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: FR 0205808
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 6
; LENGTH: 75
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-6
Query Match 100.0%; Score 110; DB 12; Length 75;
Best Local Similarity 100.0%; Pred. No. 3.6e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 PQRTKRNTRRQDVKFPG 20
Db 7 PQRTKRNTRRQDVKFPG 26

RESULT 18
US-10-431-587-4
; Sequence 4, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 4
; LENGTH: 75
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-4
Query Match 100.0%; Score 110; DB 12; Length 75;
Best Local Similarity 100.0%; Pred. No. 3.6e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 PQRTKRNTRRQDVKFPG 20
Db 7 PQRTKRNTRRQDVKFPG 26

RESULT 21
US-10-431-587-14
; Sequence 14, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 14
; LENGTH: 75
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-14
Query Match 100.0%; Score 110; DB 12; Length 75;
Best Local Similarity 100.0%; Pred. No. 3.6e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 PQRTKRNTRRQDVKFPG 20
Db 7 PQRTKRNTRRQDVKFPG 26

RESULT 19
US-10-431-587-5
; Sequence 5, Application US/10431587
; Publication No. US20040072267A1
; GENERAL INFORMATION:
; APPLICANT: BIORAD PASTEUR
; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
; FILE REFERENCE: BET 03P0456
; CURRENT APPLICATION NUMBER: US/10/431,587
; CURRENT FILING DATE: 2003-05-08
; PRIOR APPLICATION NUMBER: FR 0205808
; PRIOR FILING DATE: 2002-05-10
; NUMBER OF SEQ ID NOS: 33
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 75
; TYPE: PRT
; ORGANISM: Hepatitis C virus
US-10-431-587-5
Query Match 100.0%; Score 110; DB 12; Length 75;
Best Local Similarity 100.0%; Pred. No. 3.6e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 PQRTKRNTRRQDVKFPG 20

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Db    7 PQRKTKRNTNRRPQDVKFPG 26
      ; APPLICANT: BIORAD PASTEUR
      ; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
      ; against, an infectious microorganism
      ; FILE REFERENCE: BET 03P0456
      ; CURRENT APPLICATION NUMBER: US/10/431,587
      ; PRIORITY FILING DATE: 2003-05-08
      ; PRIORITY APPLICATION NUMBER: FR 0205808
      ; NUMBER OF SEQ ID NOS: 33
      ; SOFTWARE: PatentIn version 3.1
      ; SEQ ID NO 18
      ; LENGTH: 75
      ; TYPE: PRT
      ; ORGANISM: Hepatitis C virus
      ; FEATURE:
      ; NAME/KEY: MISC FEATURE
      ; LOCATION: (35)-(35)
      ; OTHER INFORMATION: homo-serine
      ; US-10-431-587-18

      Query Match          100.0%; Score 110; DB 12;
      Best Local Similarity 100.0%; Pred. No. 3.e-03;
      Matches 20; Conservative 0; Mismatches 0;
      Indels 0; Gaps 0;

      Qy      1 PQRKTKRNTNRRPQDVKFPG 20
      Db      7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 25
US-10-431-587-15
      ; Sequence 15, Application US/10431587
      ; Publication No. US20040072267A1
      ; GENERAL INFORMATION:
      ; APPLICANT: BIORAD PASTEUR
      ; TITLE OF INVENTION: Method for simultaneously detecting an antigen of, and an antibody
      ; against, an infectious microorganism
      ; FILE REFERENCE: BET 03P0456
      ; CURRENT APPLICATION NUMBER: US/10/431,587
      ; PRIORITY FILING DATE: 2003-05-08
      ; PRIORITY APPLICATION NUMBER: FR 0205808
      ; NUMBER OF SEQ ID NOS: 33
      ; SOFTWARE: PatentIn version 3.1
      ; SEQ ID NO 15
      ; LENGTH: 76
      ; TYPE: PRT
      ; ORGANISM: Hepatitis C virus
      ; US-10-431-587-15

      Query Match          100.0%; Score 110; DB 12;
      Best Local Similarity 100.0%; Pred. No. 3.e-03;
      Matches 20; Conservative 0; Mismatches 0;
      Indels 0; Gaps 0;

      Qy      1 PQRKTKRNTNRRPQDVKFPG 20
      Db      7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 26
US-09-578-308-1
      ; Sequence 1, Application US/09758308
      ; Patent No. US20020090607A1
      ; GENERAL INFORMATION:
      ; APPLICANT: HOWARD A. FIELDS AND YURY E. KHUDYAKOV
      ; TITLE OF INVENTION: ANTIGENIC EPITOPES AND MOSAIC POLYPEPTIDES OF HEPATITIS C VIRUS
      ; TITLE OF INVENTION: PROTEINS
      ; FILE REFERENCE: 14114.034.9U2
      ; CURRENT APPLICATION NUMBER: US/09/758,308
      ; PRIORITY FILING DATE: 2001-01-10
      ; PRIORITY APPLICATION NUMBER: 60/092,339
      ; PRIORITY FILING DATE: 1999-07-10
      ; NUMBER OF SEQ ID NOS: 5

      Query Match          100.0%; Score 110; DB 12;
      Best Local Similarity 100.0%; Pred. No. 3.e-03;
      Matches 20; Conservative 0; Mismatches 0;
      Indels 0; Gaps 0;

      Qy      1 PQRKTKRNTNRRPQDVKFPG 20
      Db      7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 24
US-10-431-587-18
      ; Sequence 18, Application US/10431587
      ; Publication No. US20040072267A1
      ; GENERAL INFORMATION:

```

SOFTWARE: PatentIn version 3.0
; SEQ ID NO: 1
; LENGTH: 91
; TYPE: PRT
; ORGANISM: Hepatitis C Virus
; US-09-756-308-1

Query Match 100.0%; Score 110; DB 9; Length 91;
Best Local Similarity 100.0%; Pred. No. 4.5e-09;
Matches 20; Conservative 0; N mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 27
US-09-756-875-8
Sequence 8, Application US/09756875
Patent No. US2002015090A1
GENERAL INFORMATION:
APPLICANT: PIKE, JAN
TITLE OF INVENTION: HEPATITIS C VIRUS PEPTIDES
NUMBER OF SEQUENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESS: Suite 701-B Columbia Square
STREET: 555 13th Street, N. W.
CITY: Washington
STATE: D. C.
COUNTRY: U. S.
ZIP: 20004

COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/756,875
FILING DATE:

CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/259,721
FILING DATE: 29-AUG-1994
APPLICATION NUMBER: PCT/GB93/00410
FILING DATE: 26-FEB-1993

ATTORNEY/AGENT INFORMATION:
NAME: ERNST, BARBARA G.
REGISTRATION NUMBER: 30,377
REFERENCE/DOCKET NUMBER: 1808-157A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 783-6040

INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 97 amino acids
TYPE: amino acid
TOPOLOGY: unknown
MOLECULE TYPE: peptide

US-09-756-875-8

Query Match 100.0%; Score 110; DB 9; Length 97;
Best Local Similarity 100.0%; Pred. No. 4.8e-09;
Matches 20; Conservative 0; N mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 7 PQRKTKRNTNRRPQDVKFPG 26

; APPLICANT: LEROUX-ROELS, Geert
; DELEYNS, Robert
; MAERTENS, Geert
; TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPEs OF HEPATITIS C VIRUS
; FILE REFERENCE: 2551-94
; CURRENT APPLICATION NUMBER: US/10/651,165
; PRIOR APPLICATION NUMBER: US/08/974,690C
; PRIOR FILING DATE: 2003-09-02
; PRIOR FILING DATE: 1997-11-19
; PRIOR APPLICATION NUMBER: PCT/EP94/03555
; PRIOR FILING DATE: 1994-10-28
; PRIOR APPLICATION NUMBER: EP 93402718.6
; PRIOR FILING DATE: 1993-11-04
; NUMBER OF SEQ ID NOS: 286
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 232
; LENGTH: 100
; TYPE: PRT
; ORGANISM: hepatitis C virus
; US-10-651-165-232

Query Match 100.0%; Score 110; DB 12; Length 100;
Best Local Similarity 100.0%; Pred. No. 4.9e-09;
Matches 20; Conservative 0; N mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 7 PQRKTKRNTNRRPQDVKFPG 26

RESULT 29
US-09-921-397-77
Sequence 77, Application US/09921397
Patent No. US2002015484A1
GENERAL INFORMATION:
APPLICANT: HYBRIGENICS
TITLE OF INVENTION: S1D nucleic acids and polypeptides selected from a pathogenic strain of the hepatitis C virus and applications thereof
FILE REFERENCE: B4809A - JAZ
CURRENT APPLICATION NUMBER: US/09/921,397
CURRENT FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: EP 00402225.7
PRIOR FILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 156
SEQ ID NO: 77
LENGTH: 103
TYPE: PRT
ORGANISM: Hepatitis C virus
; US-09-921-397-77

Query Match 100.0%; Score 110; DB 9; Length 103;
Best Local Similarity 100.0%; Pred. No. 5.1e-09;
Matches 20; Conservative 0; N mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKFPG 20
Db 20 PQRKTKRNTNRRPQDVKFPG 39

RESULT 30
US-09-851-118-14
Sequence 14, Application US/09851113B
Publication No. US20020183508A1
GENERAL INFORMATION:
APPLICANT: MAERTENS, GERT
; STUYVER, LIEVEN
; TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC AGENTS
; NUMBER OF SEQUENCES: 207

RESULT 28
US-10-651-165-232
Sequence 232, Application US/10651165
Publication No. US20040047877A1
GENERAL INFORMATION:

CORRESPONDENCE ADDRESS:
 ADDRESSEE: ARNOLD, WHITE & DURKEE
 STREET: P.O. BOX 4433
 CITY: HOUSTON
 STATE: TEXAS
 COUNTRY: USA
 ZIP: 77210-4433

COMPUTER READABLE FORM:
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/851,138
 FILING DATE: 09-May-2001
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/836,075
 FILING DATE: <Unknown>
 APPLICATION NUMBER: EP 94870166.9
 FILING DATE: 21 Oct 1994
 APPLICATION NUMBER: EP 95870076.7
 FILING DATE: 28 Jun 1995

ATTORNEY/AGENT INFORMATION:
 NAME: KAMMERER, PATRICIA A.
 REGISTRATION NUMBER: 29,775

INFORMATION FOR SEQ ID NO: 14:

SEQUENCE CHARACTERISTICS:
 LENGTH: 108 amino acids
 TYPE: amino acid
 TOPOLOGY: linear

MOLECULE TYPE: peptide
 SEQUENCE DESCRIPTION: SEQ ID NO: 14:
 US-09-851-138-14

Query Match 100.0%; Score 110; DB 9; Length 108;
 Best Local Similarity 100.0%; Pred. No. 5.3e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PORKTKRNTNRPRDVKFPG 20
 Db 7 PQRKTKRNTNRPRDVKFPG 26

RESULT 31
 US-09-321-397-78
 Sequence 78, Application US/09921397
 Patent No. US2002151484A1
 GENERAL INFORMATION:
 APPLICANT: HYBRIDGENICS
 TITLE OF INVENTION: S1D nucleic acids and polypeptides selected from a
 TITLE OF INVENTION: pathogenic strain of the hepatitis C virus and
 FILE REFERENCE: B4899A-JAZ
 CURRENT FILING DATE: 2001-08-02
 PRIOR APPLICATION NUMBER: EP 00402225.7
 NUMBER OF SEQ ID NOS: 156
 SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO 78
 LENGTH: 113
 TYPE: PRP
 ORGANISM: Hepatitis C virus
 US-09-321-397-78

Query Match 100.0%; Score 110; DB 9; Length 113;
 Best Local Similarity 100.0%; Pred. No. 5.6e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PORKTKRNTNRPRDVKFPG 20
 Db 7 PQRKTKRNTNRPRDVKFPG 26

RESULT 33
 US-10-268-569-19
 Sequence 19, Application US/10268569
 Publication No. US20030152965A1
 GENERAL INFORMATION:
 APPLICANT: Ortho-Clinical Diagnostics, Inc.
 TITLE OF INVENTION: HCV Core Protein Sequences
 FILE REFERENCE: CDS-0288
 CURRENT APPLICATION NUMBER: US/10/268,569
 CURRENT FILING DATE: 2002-10-10
 PRIOR APPLICATION NUMBER: 60/347,303
 PRIOR FILING DATE: 2001-11-11

NUMBER OF SEQ ID NOS: 19 ; Sequence 60, Application US/09851138
; SEQ ID NO: 19 ; Publication No. US2003014826781
; LENGTH: 130 ; GENERAL INFORMATION:
; TYPE: PRT ; APPLICANT: MAERTENS, GEERT
; ORGANISM: Hepatitis C virus ; TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES
; US-10-268-569-19 AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC

Query Match 100.0% Score 110; DB 14; Length 130;
Best Local Similarity 100.0% Pred. No. 6.5e-09; Indels 0; Gaps 0;
Matches 20; Conservative 0; Mismatches 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 34
US-09-851-138-46
; Sequence 46, Application US/09851138
; Publication No. US2002018508A1
; GENERAL INFORMATION:
; APPLICANT: MAERTENS, GEERT
; STUYVER, LIEVEN
; TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES
AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC
AGENTS

NUMBER OF SEQUENCES: 207
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: ARNOLD, WHITE & DURKEE
; STREET: P.O. BOX 4433
; CITY: HOUSTON
; STATE: TEXAS
; COUNTRY: USA
; ZIP: 77210-4433
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Microsoft Word 6.0 / ASCII text output

CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/851,138
; FILING DATE: 0-May-2001
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/836,075
; FILING DATE: <Unknown>
; APPLICANT NUMBER: EP 94870166.9
; FILING DATE: 21 Oct 1994
; APPLICATION NUMBER: EP 95870076.7
; FILING DATE: 28 Jun 1995
; ATTORNEY/AGENT INFORMATION:
; NAME: KAMMERER, PATRICIA A.
; REGISTRATION NUMBER: 29,775
; REFERENCE/DOCKET NUMBER: INNS:004

INFORMATION FOR SEQ ID NO: 60:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 138 amino acids
; TYPE: amino acid
; TOPOLogy: linear
; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 60:
US-09-851-138-60

Query Match 100.0% Score 110; DB 9; Length 138;
Best Local Similarity 100.0% Pred. No. 6.9e-09; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 36
US-10-292-129-14
; Sequence 14, Application US/10292129
; Publication No. US2003014826781
; GENERAL INFORMATION:
; APPLICANT: Schmidt, Emmett Vance
; APPLICANT: Chung, Raymond Taeyong
; TITLE OF INVENTION: SCRIBING ASSAY FOR HEPATITIS C VIRUS
; TITLE OF INVENTION: ANTIVIRAL AGENTS
; FILE REFERENCE: 00786-539001
; CURRENT APPLICATION NUMBER: US/10/292,129
; CURRENT FILING DATE: 2002-11-08
; PRIOR APPLICATION NUMBER: US 60/345,405
; PRIOR FILING DATE: 2001-11-09
; NUMBER OF SEQ ID NOS: 17
; SEQ ID NO 14
; LENGTH: 151
; TYPE: PRT

Query Match 100.0% Score 110; DB 9; Length 137;
Best Local Similarity 100.0% Pred. No. 6.9e-09; Mismatches 0; Indels 0; Gaps 0;
Matches 20; Conservative 0; Mismatches 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20
Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 35

; ORGANISM: Hepatitis C virus
us-10-292-14

Query Match 100.0%; Score 110; DB 14; Length 151;
Best Local Similarity 100.0%; Pred. No. 7.6e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 PQRKTKRNTNRRPQDVKEPG 20
Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 37

US-09-306-780-8

; Sequence 8, Application US/09306780

; GENERAL INFORMATION:

; APPLICANT: TAKEMURA, FUMINORI

; UENO, BIICHI

; TITLE OF INVENTION: NUCLEIC ACID-BOUND POLYPEPTIDE, METHOD

; OF PRODUCING NUCLEAR ACID-BOUND POLYPEPTIDE AND
; IMMUNOASSAY USING THE POLYPEPTIDE.

; NUMBER OF SEQUENCES: 20

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: OBLON, SPIVAK, MCCLELLAND, MAIER & NEUSTADT,

; STREET: 1755 S. JEFFERSON DAVIS HIGHWAY, SUITE 400

; CITY: ARLINGTON, VA

; STATE: VA

; COUNTRY: U.S.A.

; ZIP: 22202

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: Patent-in Release #1.0, Version #1.30

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/09/306,780

; FILING DATE: 07-MAY-1999

; CLASSIFICATION: <Unknown>

; PRIORITY NUMBER: US/08/841,657A

; ATTORNEY/AGENT INFORMATION:

; NAME: OBLON, NORMAN P.

; REGISTRATION NUMBER: 24,618

; REFERENCE/DOCKET NUMBER: 2084-033-0

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: (703) 413-3000

; TELEFAX: (703) 413-2220

; INFORMATION FOR SEQ ID NO: 8

; SEQUENCE CHARACTERISTICS:

; LENGTH: 161 amino acids

; TYPE: amino acid

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; SEQUENCE DESCRIPTION: SEQ ID NO: 8:

; US-09-306-780-8

; Query Match 100.0%; Score 110; DB 12; Length 161;

; Best Local Similarity 100.0%; Pred. No. 8.2e-09;

; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20

Db 12 PQRKTKRNTNRRPQDVKEPG 31

RESULT 38
US-09-899-046-152

; Sequence 152, Application US/09899046
; Publication No. US0030008274A1
; GENERAL INFORMATION:

; TITLE OF INVENTION: New sequences of hepatitis C virus
; TITLE OF INVENTION: genotypes for diagnosis, prophylaxis and therapy.
; NUMBER OF SEQUENCES: 270

; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent-in Release #1.0, Version #1.25 (EPO)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/362,455

; FILING DATE:

; INFORMATION FOR SEQ ID NO: 152:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/362,455

; FILING DATE:

; INFORMATION FOR SEQ ID NO: 152:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/362,455

; FILING DATE:

; INFORMATION FOR SEQ ID NO: 152:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/362,455

; FILING DATE:

; INFORMATION FOR SEQ ID NO: 152:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/362,455

; FILING DATE:

; INFORMATION FOR SEQ ID NO: 152:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/362,455

; FILING DATE:

; INFORMATION FOR SEQ ID NO: 152:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: 08/362,455

; FILING DATE:

; INFORMATION FOR SEQ ID NO: 152:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/899,046
; FILING DATE:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: 08/362,455
; FILING DATE:
; INFORMATION FOR SEQ ID NO: 152:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 166 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-899-046-152

; Query Match 100.0%; Score 110; DB 10; Length 166;
; Best Local Similarity 100.0%; Pred. No. 8.4e-09;
; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PQRKTKRNTNRRPQDVKEPG 20

Db 7 PQRKTKRNTNRRPQDVKEPG 26

RESULT 40

US-09-873-224-152
 ; Sequence 152, Application US/09873224
 ; Publication No. US2003006460A1
 ; GENERAL INFORMATION:
 ; APPLICANT: <Unknown>
 ; TITLE OF INVENTION: New sequences of hepatitis C virus
 ; NUMBER OF SEQUENCES: 270
 ; CORRESPONDENCE ADDRESS:
 ; STREET: Industriepark Zwijnaarde 7, box 4
 ; CITY: Ghent
 ; COUNTRY: Belgium
 ; ZIP: B-9052
 COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/873,224
 FILING DATE: 05-Jun-2001
 CLASSIFICATION: <Unknown>
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: 08/362,455
 ATTORNEY/AGENT INFORMATION:
 NAME: Inogenetics sa.
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 00 32 9 241 07 11
 TELEFAX: 00 32 9 241 07 99
 INFORMATION FOR SEQ ID NO: 152:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 166 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 SEQUENCE DESCRIPTION: SEQ ID NO: 152:
 US-09-873-224-152

Query Match 100.0% Score 110; DB 12; Length 166;
 Best Local Similarity 100.0%; Pred. No. 8.4e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 PORKTKRNTNRRPQDVKPG 20
 Db 7 PQRKTKRNTNRRPQDVKPG 26

Search completed: July 20, 2004, 09:45:29
 Job time : 40 secs

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OM protein - protein search, using bw model

Run on: July 20, 2004, 09:29:26 ; search time 14 Seconds
(without alignments)

Title: US-10-044-995-1

Perfect score: 106 %

Sequence: 1 MSTIPKPKRKTKRNTNRRPQ 20

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents AA:
 1: /cnr2_6.ptodata/2/iaa/5A COMB.pep:
 2: /cnr2_6.ptodata/2/iaa/5B COMB.pep:
 3: /cnr2_6.ptodata/2/iaa/6A COMB.pep:
 4: /cnr2_6.ptodata/2/iaa/6B COMB.pep:
 5: /cnr2_6.ptodata/2/iaa/POTUS COMB.pep:
 6: /cnr2_6.ptodata/2/iaa/backFiles.pep:
 *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	Match	Length	DB	ID	Description
1	106	100.0	20	2	US-08-466-975A-1		Sequence 1, Appli
2	106	100.0	20	2	US-08-391-671A-1		Sequence 1, Appli
3	106	100.0	20	3	US-08-467-902A-1		Sequence 1, Appli
4	106	100.0	20	3	US-09-275-265-1		Sequence 1, Appli
5	106	100.0	20	4	US-09-941-611-1		Sequence 1, Appli
6	106	100.0	20	4	US-09-790-497A-37		Sequence 37, Appli
7	106	100.0	20	4	US-09-790-497A-133		Sequence 133, App
8	106	100.0	20	4	US-09-576-824A-133		Sequence 133, App
9	106	100.0	22	2	US-08-146-028-37		Sequence 37, Appli
10	106	100.0	22	2	US-08-146-028-123		Sequence 123, App
11	106	100.0	22	2	US-08-146-028-124		Sequence 124, App
12	106	100.0	22	2	US-08-146-028-133		Sequence 133, App
13	106	100.0	22	3	US-08-723-425A-37		Sequence 37, Appli
14	106	100.0	22	3	US-08-723-425A-123		Sequence 123, App
15	106	100.0	22	3	US-08-723-425A-124		Sequence 124, App
16	106	100.0	22	3	US-08-723-425A-133		Sequence 133, App
17	106	100.0	22	3	US-09-112-206-37		Sequence 123, App
18	106	100.0	22	3	US-09-112-206-123		Sequence 124, App
19	106	100.0	22	3	US-09-112-206-124		Sequence 124, App
20	106	100.0	22	3	US-09-112-206-133		Sequence 133, App
21	106	100.0	22	4	US-09-576-824A-37		Sequence 37, Appli
22	106	100.0	32	4	US-09-790-497A-136		Sequence 136, App
23	106	100.0	32	4	US-09-790-497A-402		Sequence 402, App
24	106	100.0	32	4	US-09-576-824A-136		Sequence 136, App
25	106	100.0	34	4	US-09-576-824A-402		Sequence 402, App
26	106	100.0	191	2	US-08-290-665A-180		Sequence 180, App
27	106	100.0	191	5	PCT-US95-10398-180		Sequence 180, App

ALIGNMENTS

RESULT 1
US-08-466-975A-1
; Sequence 1, Application US/08466975A
; Patent No. 5910404
; GENERAL INFORMATION:
; APPLICANT: DELEYE, ROBERT J
; APPLICANT: POLLET, DIRK
; APPLICANT: MAERTENS, GEERT
; APPLICANT: VAN HEUVERS, HUGO
; TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: NIXON & VANDERHVE P.C.
; STREET: 1100 NORTH GLEBE ROAD
; CITY: ARLINGTON
; STATE: VA
; COUNTRY: USA
; ZIP: 22201
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/466,975A
; FILING DATE:
; CLASSIFICATION:
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: US/08/391,671
; FILING DATE: 13-DEC-1991
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: EP 90124241.2
; FILING DATE: 14-OCT-1992
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: WO PCT/EP91/02409
; FILING DATE: 13-DEC-1991
; PRIORITY APPLICATION DATA:
; APPLICATION NUMBER: EP 90124241.2
; FILING DATE: 14-DEC-1990
; ATTORNEY/AGENT INFORMATION:
; NAME: SADOFF, B.J.
; REGISTRATION NUMBER: 36,663
; REFERENCE/DOCKET NUMBER: 1487-5
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 703/8164000
; TELEFAX: 703/8164100
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 amino acids
; TYPE: amino acid

STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-08-466-975A-1

Query Match 100.0%; Score 106; DB 2; Length 20;
 Best Local Similarity 100.0%; Pred. No. 7e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPQRKTKRNTNRRPQ 20
 Db 1 MSTIPKPQRKTKRNTNRRPQ 20

RESULT 2
 US-08-391-671A-1
 Sequence 1, Application US/08391671A
 Patent No. 5922532
 GENERAL INFORMATION:
 APPLICANT: DELEYNS, ROBERT J
 ADDRESS: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 NUMBER OF SEQUENCES: 23
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 VIREUS
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSSEE: NIXON & VANDERHYE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/391,671
 FILING DATE: 21-FEB-1995
 CLASSIFICATION: 435
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 07/920,286
 FILING DATE: 14-OCT-1992
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 21-FEB-1995
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 13-DEC-1991
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 TELEPHONE: 7038164100
 TELEFAX: 7038164000
 REFERENCE/DOCKET NUMBER: 1487-5
 INFORMATION FOR SEQ ID NO: 1:
 LENGTH: 20 amino acids
 SEQUENCE CHARACTERISTICS:
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-08-391-671A-1

Query Match 100.0%; Score 106; DB 2; Length 20;
 Best Local Similarity 100.0%; Pred. No. 7e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPQRKTKRNTNRRPQ 20
 Db 1 MSTIPKPQRKTKRNTNRRPQ 20

RESULT 3
 US-08-467-902A-1
 Sequence 1, Application US/08467902A
 Patent No. 6007382
 GENERAL INFORMATION:
 APPLICANT: DELEYNS, ROBERT J
 ADDRESS: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/467,902A
 FILING DATE:
 CLASIFICATION:
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US/08/391,671
 FILING DATE: 14-OCT-1992
 ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 TELEPHONE: 7038164100
 TELEFAX: 7038164000
 REFERENCE/DOCKET NUMBER: 1487-5
 INFORMATION FOR SEQ ID NO: 1:
 LENGTH: 20 amino acids
 SEQUENCE CHARACTERISTICS:
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 US-08-467-902A-1

Query Match 100.0%; Score 106; DB 3; Length 20;
 Best Local Similarity 100.0%; Pred. No. 7e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPQRKTKRNTNRRPQ 20
 Db 1 MSTIPKPQRKTKRNTNRRPQ 20

RESULT 4
 US-09-275-265-1
 Sequence 1, Application US/09275265
 Patent No. 6287761
 GENERAL INFORMATION:
 APPLICANT: DELEYNS, ROBERT J

APPLICANT: POLLET, DIRK
 APPLICANT: MAERTENS, GERT
 APPLICANT: VAN HEUVERSWYN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF SEQUENCES: 23
 TITLE OF INVENTION: ANTIBODIES TO HEPATITIS C VIRUS
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTY: USA
 ZIP: 22201

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/941,611
 FILING DATE: 30-Aug-2001
 CLASSIFICATION: <Unknown>
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/391,671
 FILING DATE: 1995-02-21
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 13-DEC-1991
 APPLICATION NUMBER: EP 90124241.2
 FILING DATE: 14-DEC-1990

ATTORNEY/AGENT INFORMATION:
 NAME: SADOFF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 TELEFAX: 7038164100

INFORMATION FOR SEQ ID NO: 1:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 20 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: Peptide
 SEQUENCE DESCRIPTION: SEQ ID NO: 1:

US-09-941-611-1

Query Match 100.0%; Score 106; DB 4; Length 20;
 Best Local Similarity 100.0%; Pred. No. 7e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPQTKTKNTNRRPQ 20
 Db 1 MSTIPKPQTKTKNTNRRPQ 20

RESULT 6
 US-09-90-497A-37

GENERAL INFORMATION:
 Patent No. 6649735

PRIOR APPLICATION NUMBER: 08/790,497A
 CURRENT FILING DATE: 2001-02-23
 PRIOR APPLICATION NUMBER: 09/576,824
 PRIOR APPLICATION NUMBER: 08/723,425
 PRIOR FILING DATE: 1996-09-30
 PRIOR APPLICATION NUMBER: 09/146,028
 PRIOR FILING DATE: 1993-11-22
 PRIOR APPLICATION NUMBER: PCT/EP93/00517
 PRIOR FILING DATE: 1993-03-08
 PRIOR APPLICATION NUMBER: EP 92400598.6
 PRIOR FILING DATE: 1992-03-06
 NUMBER OF SEQ ID NOS: 600
 SOFTWARE: PatentIn Ver. 2.1

RESULT 5
 US-09-941-611-1

Sequence 1, Application US/09941611
 Patent No. 6576417

GENERAL INFORMATION:
 APPLICANT: DELEY, ROBERT J
 POLLET, DIRK
 MAERTENS, GERT
 VAN HEUVERSWYN, HUGO

TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE P.C.
 STREET: 1100 NORTH GLEBE ROAD


```

; GENERAL INFORMATION:
; APPLICANT: PROCESS FOR THE DETERMINATION OF PEPTIDES
; TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR
; TITLE OF INVENTION: BIOCUTINYLATED PEPTIDES OR BIOCUTINYLATED ANTIBODIES OR COMPOSITIONS CONTAINING THEM
; TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOCUTINYLATED EPITOPES,
; TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
; TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
; NUMBER OF SEQUENCES: 453
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/146,028
; INFORMATION FOR SEQ ID NO: 123;
; SEQUENCE CHARACTERISTICS:
; LENGTH: 22 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; INDIVIDUAL ISOLATE: HCV
; FEATURE:
; NAME/KEY: Xaa is H2N
; LOCATION: 1
; FEATURE:
; NAME/KEY: Xaa is CONH2
; LOCATION: 22
; US-08-146-028-123

Query Match      100.0%; Score 106; DB 2; Length 22;
Best Local Similarity 100.0%; Pred. No. 7.7e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy   1 MSTIPKPKRQTKYKNTNRRPQ 20
     ||||| | | | | | | | | | |
Db   2 MSTIPKPKRQTKYKNTNRRPQ 21

; RESULT 11
; US-08-146-028-124
; Sequence 124, Application US/08146028
; Parent No. 5891640
; GENERAL INFORMATION:
; APPLICANT: PROCESS FOR THE DETERMINATION OF PEPTIDES
; TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR
; TITLE OF INVENTION: BIOCUTINYLATED PEPTIDES OR BIOCUTINYLATED ANTIBODIES OR COMPOSITIONS CONTAINING THEM
; TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOCUTINYLATED EPITOPES,
; TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPES,
; TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM
; NUMBER OF SEQUENCES: 453
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/146,028
; INFORMATION FOR SEQ ID NO: 124;
; SEQUENCE CHARACTERISTICS:
; LENGTH: 22 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; HYPOTHETICAL: NO
; ORIGINAL SOURCE:
; INDIVIDUAL ISOLATE: HCV
; FEATURE:
; NAME/KEY: Xaa is H2N
; LOCATION: 1

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STATE: VA
COUNTRY: USA
ZIP: 22201
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/723, 425A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 1487-13
TELEPHONE: 703-816-4000
TELEFAX: 703-816-4100
INFORMATION FOR SEQ ID NO: 123:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: Peptide
HYPOTHETICAL: NO
ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: HCV
FEATURE:
NAME/KEY: Xaa is H2N
LOCATION: 1
FEATURE:
NAME/KEY: Xaa is CONH2
LOCATION: 22
US-08-723-425A-123

Query Match 100.0% Score 106; DB 3; Length 22;
Best Local Similarity 100.0%; Pred. No. 7.e-09;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIIPKPKRTKNTNRRPQ 20
Db 2 MSTIIPKPKRTKNTNRRPQ 21

RESULT 15
US-08-723-425A-124
Sequence 124, Application US/08723425A
Patent No. 6165730
GENERAL INFORMATION:
APPLICANT: DELEYS, ROBERT
TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF
PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
TITLE OF INVENTION: PEPTIDES AND THEIR USE IN A PROCESS FOR DETERMINATION OF
TITLE OF INVENTION: ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING ...
NUMBER OF SEQUENCES: 453
CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE, P.C.
STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
CITY: Arlington
STATE: VA
COUNTRY: USA
ZIP: 22201
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/723, 425A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 1487-13
TELEPHONE: 703-816-4000
TELEFAX: 703-816-4100
INFORMATION FOR SEQ ID NO: 124:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: Peptide
HYPOTHETICAL: NO

ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: HCV
FEATURE:
NAME/KEY: Xaa is H2N
LOCATION: 1

FEATURE:
NAME/KEY: Xaa is Gly-Gly-Lys (Bio) -CONH2
LOCATION: 22

US-08-723-425A-124

Query Match 16
Best Local Similarity 100.0%; Score 106; DB 3; Length 22;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
Db 2 MSTIPKPKRKTKRNTNRRPQ 21

RESULT 17
US-09-112-206-37
; Sequence 37, Application US/09112206

GENERAL INFORMATION:
APPLICANT: ; TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPIES AND THEIR LENGTH: 22; ; PATENT NO. 6210903

APPLICANT: ; TITLE OF INVENTION: PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPIES, 1 ; TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPIES, 1 ; NUMBER OF SEQUENCES: 453

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (BPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/112,206

FILING DATE:
PRIORITY APPLICATION DATA:
APPLICATION NUMBER: US 08/146,028
FILING DATE:
PRIORITY NUMBER: US 08/146,028
INFORMATION FOR SEQ ID NO: 37:

SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO

INDIVIDUAL ISOLATE: HCV
FEATURE:
NAME/KEY: Modified-site
LOCATION: 1
FEATURE:
NAME/KEY: Modified-site
LOCATION: 22

US-09-112-206-37

Query Match 18
Best Local Similarity 100.0%; Score 106; DB 3; Length 22;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
Db 2 MSTIPKPKRKTKRNTNRRPQ 21

RESULT 18
US-09-112-206-123
; Sequence 123, Application US/09112206

GENERAL INFORMATION:
APPLICANT: ; TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPIES AND THEIR LENGTH: 22; ; PATENT NO. 6210903

APPLICANT: ; TITLE OF INVENTION: PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPIES, 1 ; TITLE OF INVENTION: PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPIES, 1 ; NUMBER OF SEQUENCES: 453

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (BPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/723,425A

FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: SADOFF, B.J.
REGISTRATION NUMBER: 36,663
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703-816-4100
INFORMATION FOR SEQ ID NO: 133:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO

ORIGINAL SOURCE:
INDIVIDUAL ISOLATE: HCV
FEATURE:
NAME/KEY: Xaa is absent
LOCATION: 1

FEATURE:
NAME/KEY: Xaa is absent
LOCATION: 22

US-08-723-425A-133

Query Match 19
Best Local Similarity 100.0%; Score 106; DB 3; Length 22;
Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

APPLICATION NUMBER: US/09/112,206
 FILING DATE:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/146,028
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 123:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 22 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Xaa is H2N
 LOCATION: 1
 FEATURE:
 NAME/KEY: Xaa is CONH2
 LOCATION: 22
 US-09-112-206-123
 Query Match 100.0%; Score 106; DB 3; Length 22;
 Best Local Similarity 100.0%; Pred. No. 7.e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Qy 1 MSTIPKQPKTKRNTNRRPQ 20
 Db 2 MSTIPKQPKTKRNTNRRPQ 21

RESULT 19
 US-09-112-206-124
 Sequence 124, Application US/091122206
 Patent No. 6210903
 GENERAL INFORMATION:
 APPLICANT:
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM NUMBER OF SEQUENCES: 453
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/112,206
 FILING DATE:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/146,028
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 124:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 22 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Xaa is H2N
 LOCATION: 1
 FEATURE:
 NAME/KEY: Xaa is Gly-Gly-Lys (Bio) -CONH2
 LOCATION: 22
 US-09-112-206-124
 Query Match 100.0%; Score 106; DB 3; Length 22;
 Best Local Similarity 100.0%; Pred. No. 7.e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

RESULT 20
 US-09-112-206-133
 Sequence 133, Application US/09112206
 Patent No. 6210903
 GENERAL INFORMATION:
 APPLICANT:
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: IN A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM NUMBER OF SEQUENCES: 453
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/112,206
 FILING DATE:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 08/146,028
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 125:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 22 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Xaa is H2N
 LOCATION: 1
 FEATURE:
 NAME/KEY: Xaa is Gly-Gly-Lys (Bio) -CONH2
 LOCATION: 22
 US-09-112-206-133
 Query Match 100.0%; Score 106; DB 3; Length 22;
 Best Local Similarity 100.0%; Pred. No. 7.e-09;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

RESULT 21
 US-09-576-824A-37
 Sequence 37, Application US/09576624A
 Patent No. 6667387
 GENERAL INFORMATION:
 APPLICANT: De Leyts, Robert
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: A PROCESS FOR DETERMINATION OF ANTIBODIES OR BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR LENGTH: 22
 TITLE OF INVENTION: EPITOPE, A PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM NUMBER OF SEQUENCES: 453
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US 08/146,028
 FILING DATE:
 INFORMATION FOR SEQ ID NO: 126:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 22 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ORIGINAL SOURCE:
 INDIVIDUAL ISOLATE: HCV
 FEATURE:
 NAME/KEY: Xaa is H2N
 LOCATION: 1
 FEATURE:
 NAME/KEY: Xaa is Gly-Gly-Lys (Bio) -CONH2
 LOCATION: 22
 US-09-576-824A-37
 Query Match 100.0%; Score 106; DB 3; Length 22;
 Best Local Similarity 100.0%; Pred. No. 7.e-09;

PRIOR APPLICATION NUMBER: 09/146,028
 PRIOR FILING DATE: 1993-11-22
 PRIOR APPLICATION NUMBER: PCT/EP93/00517
 PRIOR FILING DATE: 1993-03-08
 PRIOR APPLICATION NUMBER: EP 92400598.6
 PRIOR FILING DATE: 1992-03-06
 NUMBER OF SEQ ID NOS: 600
 SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO 37
 LENGTH: 22
 TYPE: PRT
 ORGANISM: Hepatitis C virus
 FEATURE:
 NAME/KEY: VARIANT
 LOCATION: (1)
 OTHER INFORMATION: modified site
 NAME/KEY: VARIANT
 LOCATION: (22)
 OTHER INFORMATION: modified site
 US-09-576-824A-37

Query Match 100.0% Score 106; DB 4; Length 22;
 Best Local Similarity 100.0%; Pred. No. 7.e-0; Mismatches 0; Indels 0; Gaps 0;
 Matches 20; Conservative 0;
 Qy 1 MSTIPKPORTKTKNTNRRPQ 20
 Db 2 MSTIPKPORTKTKNTNRRPQ 21

RESULT 22

US-09-790-497A-136

Sequence 136, Application US/09790497A

Patent No. 6649735

GENERAL INFORMATION:

APPLICANT: De Leyts,

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING

TO IMMUNOLOGICALLY IMPORTANT EPITOPES AND THEIR USE IN

A PROCESS FOR DETERMINATION OF ANTIBODIES OF

BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT

TITLE OF INVENTION: EPITOPEs, A PROCESS FOR PREPARING THEM AND COMPOSITIONS

TITLE OF INVENTION: CONTAINING THEM

FILE REFERENCE: 2752-16

CURRENT APPLICATION NUMBER: US/09/790,497A

CURRENT FILING DATE: 2001-02-23

PRIOR APPLICATION NUMBER: 09/576,84

PRIOR FILING DATE: 2000-05-23

PRIOR APPLICATION NUMBER: 09/723,425

PRIOR FILING DATE: 1996-09-30

PRIOR APPLICATION NUMBER: 09/146,028

PRIOR FILING DATE: 1993-11-22

PRIOR FILING DATE: 1993-03-08

NUMBER OF SEQ ID NOS: 600

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 136

LENGTH: 32

TYPE: PRT

ORGANISM: Hepatitis C virus

US-09-790-497A-136

Query Match 100.0% Score 106; DB 4; Length 32;
 Best Local Similarity 100.0%; Pred. No. 1.e-0; Mismatches 0; Indels 0; Gaps 0;
 Matches 20; Conservative 0;

Qy 1 MSTIPKPORTKTKNTNRRPQ 20

Db 1 MSTIPKPORTKTKNTNRRPQ 20

RESULT 23

Query Match 100.0% Score 106; DB 4; Length 32;

US-09-576-824A-136

Query Match 100.0% Score 106; DB 4; Length 32;

US-09-576-824A-136

RESULT 23

US-09-790-497A-402
 Sequence 402, Application US/09790497A
 Patent No. 6649735
 GENERAL INFORMATION: Robert
 APPLICANT: De Leyts,
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
 TO IMMUNOLOGICALLY IMPORTANT EPITOPEs AND THEIR USE IN
 A PROCESS FOR DETERMINATION OF ANTIBODIES OF
 BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
 TITLE OF INVENTION: EPITOPEs, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
 TITLE OF INVENTION: CONTAINING THEM

CURRENT APPLICATION NUMBER: US/09/790,497A
 CURRENT FILING DATE: 2001-02-23
 PRIOR APPLICATION NUMBER: 09/576,824
 PRIOR FILING DATE: 2000-05-23
 PRIOR APPLICATION NUMBER: 09/723,425
 PRIOR FILING DATE: 1996-09-30
 PRIOR APPLICATION NUMBER: 09/146,028
 PRIOR FILING DATE: 1993-11-22
 PRIOR APPLICATION NUMBER: PCT/EP93/00517
 PRIOR FILING DATE: 1993-03-08
 NUMBER OF SEQ ID NOS: 600
 SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO 402
 LENGTH: 32
 TYPE: PRT
 ORGANISM: Hepatitis C virus

US-09-790-497A-402
 Sequence 402, Application US/09790497A
 Patent No. 6649735
 GENERAL INFORMATION: Robert
 APPLICANT: De Leyts,
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
 TO IMMUNOLOGICALLY IMPORTANT EPITOPEs AND THEIR USE IN
 A PROCESS FOR DETERMINATION OF ANTIBODIES OF
 BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
 TITLE OF INVENTION: EPITOPEs, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
 TITLE OF INVENTION: CONTAINING THEM

Query Match 100.0% Score 106; DB 4; Length 32;
 Best Local Similarity 100.0%; Pred. No. 1.e-0; Mismatches 0; Indels 0; Gaps 0;
 Matches 20; Conservative 0;

Qy 1 MSTIPKPORTKTKNTNRRPQ 20
 Db 1 MSTIPKPORTKTKNTNRRPQ 20

RESULT 24
 US-09-576-824A-136
 Sequence 136, Application US/09576824A
 Patent No. 6667387
 GENERAL INFORMATION: Robert
 APPLICANT: De Leyts,
 TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING
 TO IMMUNOLOGICALLY IMPORTANT EPITOPEs AND THEIR USE IN
 A PROCESS FOR DETERMINATION OF ANTIBODIES OF
 BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT
 TITLE OF INVENTION: EPITOPEs, A PROCESS FOR PREPARING THEM AND COMPOSITIONS
 TITLE OF INVENTION: CONTAINING THEM

CURRENT APPLICATION NUMBER: US/09/576,824A
 CURRENT FILING DATE: 2000-05-23
 PRIOR APPLICATION NUMBER: 09/576,824
 PRIOR FILING DATE: 1996-09-30
 PRIOR APPLICATION NUMBER: 09/146,028
 PRIOR FILING DATE: 1993-11-22
 PRIOR APPLICATION NUMBER: PCT/EP93/00517
 PRIOR FILING DATE: 1993-03-08
 NUMBER OF SEQ ID NOS: 600
 SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO 136
 LENGTH: 32
 TYPE: PRT
 ORGANISM: Hepatitis C virus

US-09-576-824A-136

Best Local Similarity 100.0%; Pred. No. 1.1e-08; Mismatches 0; Indels 0; Gaps 0;

Matches 20; Conservative 0; Nucleotides 0;

US-09-576-824A-402 Sequence 402, Application US/09576824A

Patent No. 6667387

GENERAL INFORMATION:

APPLICANT: De Ley's, Robert

TITLE OF INVENTION: PROCESS FOR THE DETERMINATION OF PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE AND THEIR USE IN A PROCESS FOR DETERMINATION OF ANTIBODIES OF BIOTINYLATED PEPTIDES CORRESPONDING TO IMMUNOLOGICALLY IMPORTANT EPITOPE, A PROCESS FOR PREPARING THEM AND COMPOSITIONS CONTAINING THEM

FILE REFERENCE: 2752-11

CURRENT APPLICATION NUMBER: US/09/576,824A

PRIOR APPLICATION NUMBER: 08/7023,425

PRIOR FILING DATE: 1996-09-30

PRIOR APPLICATION NUMBER: 09/146,028

PRIOR FILING DATE: 1993-11-22

PRIOR APPLICATION NUMBER: PCT/EP93/00517

PRIOR FILING DATE: 1993-03-08

PRIOR APPLICATION NUMBER: EP 92400598.6

PRIOR FILING DATE: 1992-03-06

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO: 402 LENGTH: 34

TYPE: PCT ORGANISM: Hepatitis C virus FEATURE: NAME/KEY: VARIANT LOCATION: (1)

OTHER INFORMATION: Xaa = modified site : when present, represents an amino acid, amino group, or chemically modified amino terminus

NAME/KEY: VARIANT LOCATION: (34)

OTHER INFORMATION: Xaa = modified site : when present, represents an amino acid, OH-group, NH2-group, or a linkage involving these two groups

US-09-576-824A-402

Query Match Score 100%; Length 34; Best Local Similarity 100.0%; Pred. No. 1.2e-08; Mismatches 0; Indels 0; Gaps 0;

Matches 20; Conservative 0; Nucleotides 0;

US-09-576-824A-402

Query Match Score 100%; Length 34; Best Local Similarity 100.0%; Pred. No. 1.2e-08; Mismatches 0; Indels 0; Gaps 0;

Matches 20; Conservative 0; Nucleotides 0;

US-08-290-665A-180 Sequence 180, Application US/08290665A

Patent No. 5882852

GENERAL INFORMATION:

APPLICANT: BURKH, J., MILLER, R. H. AND PURCELL, R. H.

TITLE OF INVENTION: NUCLEOTIDE AND DEDUCED AMINO ACID SEQUENCES OF THE ENVELOPE 1 AND CORE GENES OF ISOLATES OF HEPATITIS C VIRUS

TITLE OF INVENTION: AMINO ACID SEQUENCES OF THE ENVELOPE 1 AND CORE GENES OF ISOLATES OF HEPATITIS C VIRUS

TITLE OF INVENTION: AND THE USE OF REAGENTS DERIVED FROM THESE SEQUENCES IN DIAGNOSTIC METHODS AND VACCINES

NUMBER OF SEQUENCES: 263

CORRESPONDENCE ADDRESS:

ADDRESSEE: MORGAN & FINNEGAN

STREET: 345 PARK AVENUE

CITY: NEW YORK

STATE: NEW YORK

COUNTRY: USA

ZIP: 10154

COMPUTER READABLE FORM:

MEDIUM TYPE: FLOPPY DISK

COMPUTER: IBM PC COMPATIBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: WORDPERFECT 5.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: PCT/US95/10398

FILING DATE: 15-AUG-1995

CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/086,428

FILING DATE: 29 JUNE 1993
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: 08/200/665
 FILING DATE: 15 AUGUST 1994
 NAME: RICHARD W. BORK
 REGISTRATION NUMBER: 36,459
 REFERENCE/DOCKET NUMBER: 2026-4116
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (212) 758-4800
 TELEFAX: (212) 751-6849
 TELEX: 421792
 INFORMATION FOR SEQ ID NO: 180:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 191 amino acids
 STRANDEDNESS: unknown
 TOPOLOGY: unknown
 ORGANISM: *homo sapiens*
 INDIVIDUAL ISOLATE: T2
 PCR-US95-10398-180

Query Match Score 106; DB 5; Length 191;
 Best Local Similarity 100.0%; Pred. No. 6.e-0;
 Matches 20; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;

Qy 1 MSTIPKPORTRKTRNRRPQ 20
 Db 1 MSTIPKPORTRKTRNRRPQ 20

RESULT 28
 US-08-635-886C-181
 Sequence 181, Application US/08635886C
 Patent No. 6555114
 GENERAL INFORMATION:
 APPLICANT: LEROUX-ROELS, Geert
 APPLICANT: DELEY, Robert
 APPLICANT: MAERTENS, Geert
 TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C
 FILE REFERENCE: 2752-18
 CURRENT APPLICATION NUMBER: US/08/635,886C
 CURRENT FILING DATE: 1996-04-25
 PRIOR APPLICATION NUMBER: PCT/EP94/03555
 PRIOR FILING DATE: 1994-10-28
 PRIOR APPLICATION NUMBER: EP 93402718.6
 PRIOR FILING DATE: 1993-11-04
 NUMBER OF SEQ ID NOS: 286
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO 181
 LENGTH: 450
 TYPE: PRT
 ORGANISM: hepatitis C virus
 US-08-635-886C-181

Query Match Score 106; DB 4; Length 450;
 Best Local Similarity 100.0%; Pred. No. 1.5e-0;
 Matches 20; Conservative 0; Mismatches 0;
 Indels 0; Gaps 0;

Qy 1 MSTIPKPORTRKTRNRRPQ 20
 Db 1 MSTIPKPORTRKTRNRRPQ 20

RESULT 29
 US-08-974-690C-181
 Sequence 181, Application US/08974690C
 Patent No. 661333
 GENERAL INFORMATION:
 APPLICANT: LEROUX-ROELS, Geert
 APPLICANT: DELEYS, Robert

APPLICANT: MAERTENS, Geert
 APPLICANT: VAN HEUVERSIN, HUGO
 APPLICANT: DELEY, Robert
 APPLICANT: POLLET, DIRK
 APPLICANT: MAERTENS, Geert
 APPLICANT: VAN HEUVERSIN, HUGO
 TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF
 ANTIBODIES TO HEPATITIS C VIRUS
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: NIXON & VANDERHVE P.C.
 STREET: 1100 NORTH GLEBE ROAD
 CITY: ARLINGTON
 STATE: VA
 COUNTRY: USA
 ZIP: 22201
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.3.0
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/466,975A
 FILING DATE:
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US/08/391,671
 FILING DATE:
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: US 07/920,286
 FILING DATE: 14-OCT-1992
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: WO PCT/EP91/02409
 FILING DATE: 13-DEC-1991
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: EP 90124241-2
 FILING DATE: 14-DEC-1990
 ATTORNEY/AGENT INFORMATION:
 NAME: SADEF, B.J.
 REGISTRATION NUMBER: 36,663
 REFERENCE/DOCKET NUMBER: 1487-5
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 7038164000
 TELEFAX: 7038164100

INFORMATION FOR SEQ ID NO: 23:

SEQUENCE CHARACTERISTICS:
 LENGTH: 2894 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: peptide
 HYPOTHETICAL: NO
 ANTI-SENSE: NO

US-08-466-975A-23.

Query Match

Best Local Similarity 100.0%; Pred. No. 9.8e-07; Length 2894; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20

Db 1 MSTIPKPKRKTKRNTNRRPQ 20

RESULT 31

Sequence 23, Application US/08391671A

Patent No. 5922532

GENERAL INFORMATION:

APPLICANT: DELEYS, ROBERT J

APPLICANT: POLLET, DIRK

APPLICANT: MAERTENS, GERT

APPLICANT: VAN HEUVERWYN, HUGO

TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF

TITLE OF INVENTION: ANTIBODIES TO HEPATITIS C VIRUS

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:

ADDRESSEE: NIXON & VANDERHYE P.C.

STREET: 1100 NORTH GLEBE ROAD

CITY: ARLINGTON

STATE: VA

COUNTRY: USA

ZIP: 22201

COMPUTER READABLE FORM:

APPLICATION NUMBER: US/08/391,671

FILING DATE: 21-FEB-1995

CLASSIFICATION: 435

PRIORITY APPLICATION DATA:

APPLICATION NUMBER: EP 90124241.2

FILING DATE: 14-OCT-1992

PRIORITY APPLICATION DATA:

APPLICATION NUMBER: WO PCT/EP91/02409

FILING DATE: 13-DEC-1991

PRIORITY APPLICATION DATA:

APPLICATION NUMBER: EP 90124241.2

FILING DATE: 14-DEC-1990

ATTORNEY/AGENT INFORMATION:

NAME: SADOFF, B.J.

REGISTRATION NUMBER: 36,663

REFERENCE/DOCKET NUMBER: 1487-5

TELECOMMUNICATION INFORMATION:

TELEPHONE: 7038164000

TELEFAX: 7038164100

INFORMATION FOR SEQ ID NO: 23:

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

INFORMATION FOR SEQ ID NO: 23:

SEQUENCE CHARACTERISTICS:

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-391-671A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-466-975A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-466-975A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-08-467-902A-23

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

Db 1 MSTIPKPKRTKTKNTNRRPQ 20

RESULT 33

US-09-275-265-23

; Sequence 23; Application US/09275265

; Patent No. 6287761

; GENERAL INFORMATION:

; APPLICANT: DELEY, ROBERT J

; APPLICANT: POLLET, DIRK

; APPLICANT: MAERTENS, GEERT

; APPLICANT: VAN HEUVERSWYN, HUGO

; TITLE OF INVENTION: SYNTHETIC ANTIGENS FOR THE DETECTION OF ANTIBODIES TO HEPATITIS C VIRUS

; NUMBER OF SEQUENCES: 23

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: NIXON & VANDERHVE P.C.

; STREET: 1100 NORTH GLEBE ROAD

; CITY: ARLINGTON

; STATE: VA

; COUNTRY: USA

; ZIP: 22201

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/275,265

FILING DATE:

CLASSIFICATION:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/06/391,671

FILING DATE: 21-FEB-1995

APPLICATION NUMBER: US 07/920,286

FILING DATE: 14-OCT-1992

PRIOR APPLICATION DATA:

APPLICATION NUMBER: WO PCT/EP91/02409

FILING DATE: 13-DEC-1991

PRIOR APPLICATION DATA:

APPLICATION NUMBER: EP 90124241.2

FILING DATE: 14-DEC-1991

ATTORNEY/AGENT INFORMATION:

NAME: SADOFF, B.J.

REGISTRATION NUMBER: 36,663

REFERENCE/DOCKET NUMBER: 1487-5

TELECOMMUNICATION INFORMATION:

TELEPHONE: 7038164000

TELEFAX: 7038164100

INFORMATION FOR SEQ ID NO: 23:

SEQUENCE CHARACTERISTICS:

LENGTH: 2894 amino acids

TYPE: amino acid

STRANDBNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

ANTI-SENSE: NO

US-09-275-265-23

Query Match 100.0%; Score 106; DB 3; Length 2894;

Best Local Similarity 100.0%; Pred. No. 9.8e-07; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRTKTKNTNRRPQ 20

Db 1 MSTIPKPKRTKTKNTNRRPQ 20

RESULT 34

US-09-611-23

; Sequence 23; Application US/09941611

Query Match 100.0%; Score 106; DB 3; Length 2894;

Best Local Similarity 100.0%; Pred. No. 9.8e-07; Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRTKTKNTNRRPQ 20

Db 1 MSTIPKPKRTKTKNTNRRPQ 20

RESULT 35

US-08-635-886C-232

; Sequence 232; Application US/08635886C

; Patent No. 655114

; GENERAL INFORMATION:

; APPLICANT: LEROUX-ROELS, Geert

; APPLICANT: DELEY, Robert

; APPLICANT: MAERTENS, Geert

; TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOSES OF HEPATITIS C VIRUS

; FILE REFERENCE: 2752-18

; CURRENT APPLICATION NUMBER: US/08/635,886C

; CURRENT FILING DATE: 1996-04-25

PRIOR APPLICATION NUMBER: PCT/EP94/03555
 PRIOR FILING DATE: 1994-10-28
 PRIOR APPLICATION NUMBER: EP 93402718.6
 PRIOR FILING DATE: 1993-11-04
 SEQ ID NO: 232
 SOFTWARE: PatentIn version 3.1
 LENGTH: 100
 TYPE: PRT
 ORGANISM: hepatitis C virus
 US-08-635-886C-232

Query Match 98.1%; Score 104; DB 4; Length 100;
 Best Local Similarity 95.0%; Pred. No. 6.6e-08;
 Matches 19; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
 Db 1 MSTLPKPKRKTKRNTNRRPQ 20

RESULT 36
 US-08-974-690C-232
 Sequence 232, Application US/08974690C
 Patent No. 6613333
 GENERAL INFORMATION:
 APPLICANT: LEROIX-ROELIS, Geert
 APPLICANT: DELEYNS, Robert
 APPLICANT: MAERTENS, Geert
 TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPE OF HEPATITIS C VIRUS
 FILE REFERENCE: 2551-94
 CURRENT FILING DATE: 1997-11-19
 PRIOR APPLICATION NUMBER: PCT/EP94/03555
 PRIOR FILING DATE: 1994-10-28
 PRIOR APPLICATION NUMBER: EP 93402718.6
 PRIOR FILING DATE: 1993-11-04
 NUMBER OF SEQ ID NOS: 286
 SOFTWARE: PatentIn version 3.1
 SEQ ID NO: 232
 LENGTH: 100
 TYPE: PRT
 ORGANISM: hepatitis C virus
 US-08-974-690C-232

Query Match 98.1%; Score 104; DB 4; Length 100;
 Best Local Similarity 95.0%; Pred. No. 6.6e-08;
 Matches 19; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
 Db 1 MSTLPKPKRKTKRNTNRRPQ 20

RESULT 37
 US-08-836-075A-46
 Sequence 46, Application US/08836075A
 Patent No. 6180758
 GENERAL INFORMATION:
 APPLICANT: MAERTENS, GEERT
 APPLICANT: STUYVER, LIEVEN
 TITLE OF INVENTION: NEW SEQUENCES OF HEPATITIS C VIRUS GENOTYPES
 NUMBER OF SEQUENCES: 207
 TITLE OF INVENTION: AGENTS AND THEIR USE AS PROPHYLACTIC, THERAPEUTIC AND DIAGNOSTIC
 ADDRESSEE: ARNOLD, WHITE & DURKEE
 STREET: P.O. BOX 4433
 CITY: HOUSTON
 STATE: TEXAS
 COUNTRY: USA
 ZIP: 77210-4433

COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC Compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Microsoft Word 6.0 / ASCII text output
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/836,075A
 FILING DATE: 21 Apr 1997
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: PCT/BP95/04155
 FILING DATE: 23 Oct 1995
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 94870166.9
 FILING DATE: 21 Oct 1994
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: EP 95870076.7
 FILING DATE: 28 Jun 1995
 ATTORNEY/AGENT INFORMATION:
 NAME: KAMMERER, PATRICIA A.
 REGISTRATION NUMBER: 29,775
 REFERENCE/DOCKET NUMBER: INNS: 004
 INFORMATION FOR SEQ ID NO: 46:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 137 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: Peptide
 US-08-836-075A-46

Query Match 98.1%; Score 104; DB 3; Length 137;
 Best Local Similarity 95.0%; Pred. No. 9.1e-08;
 Matches 19; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MSTIPKPKRKTKRNTNRRPQ 20
 Db 1 MSTLPKPKRKTKRNTNRRPQ 20

RESULT 38
 US-07-946-054-9
 Sequence 9, Application US/07946054
 Patent No. 5582368
 GENERAL INFORMATION:
 APPLICANT: Wang, Chang Yi
 APPLICANT: Hosein, Barbara H
 TITLE OF INVENTION: No. 5582961el Branched Hybrid and Cluster
 TITLE OF INVENTION: No. 5582961el Branched Hybrid and Cluster
 TITLE OF INVENTION: Peptides Effective in Diagnosing and Detecting No. 5582968-A,
 NUMBER OF SEQUENCES: 12
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: United Biomedical Inc.
 STREET: 25 Davids Dr.
 CITY: Hauppauge
 STATE: New York
 COUNTRY: USA
 ZIP: 11788
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC Compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/07/946,054
 FILING DATE: 15-SEP-1992
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: Wilson, M. Lisa
 REGISTRATION NUMBER: 34,045
 REFERENCE/DOCKET NUMBER: 2000
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 516-273-2828
 TELEFAX: 516-273-1717
 INFORMATION FOR SEQ ID NO: 9:

SEQUENCE CHARACTERISTICS:
; LENGTH: 61 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide

US-07-946-054-9

Query Match 95.3%; Score 101; DB 1; Length 61;
Best Local Similarity 100.0%; Prod. No. 1.1e-07;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 STIPKPKRTKRTNRRPQ 20
Db 1 STIPKPKRTKRTNRRPQ 19

RESULT 39

US-08-083-947-23 Application US/08083947
; Patent No. 5639594
; GENERAL INFORMATION:
; APPLICANT: Wang, Chang Yi
; TITLE OF INVENTION: No. 5639594-1 Linear and Branched Peptides Effective
; TITLE OF INVENTION: in Diagnosing and Detecting No. 5639594-A, No. 5639594-B Hepat
; NUMBER OF SEQUENCES: 23
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: M. Lisa Wilson
; STREET: 25 Davids Drive
; CITY: Hauppauge
; STATE: NY
; COUNTRY: USA
; ZIP: 11788
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/530,550
; FILING DATE:
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: Wilson, M. Lisa
; REGISTRATION NUMBER: 34,045
; REFERENCE/DOCKET NUMBER: 2000Z
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (516)273-2828
; TELEFAX: (516)273-1717
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 61 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide

US-08-530-550-3

Query Match 95.3%; Score 101; DB 1; Length 61;
Best Local Similarity 100.0%; Prod. No. 1.1e-07;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 STIPKPKRTKRTNRRPQ 20
Db 1 STIPKPKRTKRTNRRPQ 19

Search completed: July 20, 2004, 09:37:15
Job time : 14 secs

US-08-083-947-23

Query Match 95.3%; Score 101; DB 1; Length 61;
Best Local Similarity 100.0%; Prod. No. 1.1e-07;
Matches 19; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 STIPKPKRTKRTNRRPQ 20
Db 1 STIPKPKRTKRTNRRPQ 19

RESULT 40

US-08-530-550-3
; Sequence 3, Application US/08530550
; Patent No. 5736321

